OM protein - protein search, using sw model

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GenCore version 5.1.6

Run on: December 4, 2003, 14:31:50 ; Search time 35.2613 Seconds
              (without alignments)
              741.553 Million cell updates/sec

Title: US-08-569-749-2
Perfect Score: 3277
Sequence: 1 MHKTASQRLFPGPSYQNIKS.........LRKCP:CRGIIKGTVRTPLS 618

Scoring table: BLOSUM62
              Gapop 10.0 , Gapext 0.5

Searched: 328717 seqs, 42310858 residues

Total number of hits satisfying chosen parameters: 328717

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   Issued_Patents_AA:*
   1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
   2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
   3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
   4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
   5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
   6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	3277	100.0	618	3	US-08-569-749-2	Sequence 2, Appli
2	3277	100.0	618	4	US-09-069-023-29	Sequence 29, Appl
3	3277	100.0	618	2	PCT-US96-12860-2	Sequence 2, Appli
4	3247	99.1	618	2	US-09-212-971-8	Sequence 8, Appli
5	3247	99.1	618	3	US-08-511-485-8	Sequence 8, Appli
6	3247	99.1	618	3	US-08-800-929A-8	Sequence 8, Appli
7	3247	99.1	618	4	US-09-617-053A-8	Sequence 8, Appli
8	3247	99.1	618	4	US-09-201-936-8	Sequence 8, Appli
9	3247	99.1	618	3	US-08-569-749-14	Sequence 14, Appl
10	2728	83.2	612	5	PCT-US96-12860-14	Sequence 14, Appl
11	2724	83.1	612	5	US-09-212-971-14	Sequence 14, Appl
12	2724	83.1	612	3	US-08-800-929A-14	Sequence 14, Appl
13	2724	83.1	612	4	US-09-617-053A-14	Sequence 14, Appl
14	2654	81.0	591	4	US-09-201-936-42	Sequence 42, Appl
15	2353	71.8	604	3	US-08-569-749-4	Sequence 4, Appl
16	2353	71.8	604	2	PCT-US96-12860-4	Sequence 4, Appl
17	2332	71.2	604	5	US-08-511-485-6	Sequence 6, Appl
18	2332	71.2	604	3	US-09-212-971-6	Sequence 6, Appl
19	2332	71.2	604	4	US-08-800-929A-6	Sequence 6, Appl
20	2332	71.2	604	4	US-09-617-053A-6	Sequence 6, Appl
21	2332	71.2	604	4	US-09-201-936-6	Sequence 6, Appli
22	2326	71.0	438	5	PCT-US95-05922A-2	Sequence 2, Appli
23	2172	66.3	600	3	US-09-212-971-12	Sequence 12, Appli
24	2172	66.3	600	3	US-08-800-929A-12	Sequence 12, Appl
25	2172	66.3	600	4	US-09-617-053A-12	Sequence 12, Appl
26	2152	65.7	602	4	US-09-201-936-40	Sequence 40, Appl
27	909	27.7	497	2	US-08-511-485-4	Sequence 4, Appl
28	909	27.7	497	3	US-09-212-971-4	Sequence 4, Appli
29	909	27.7	497	3	US-08-800-929A-4	Sequence 4, Appli
30	909	27.7	497	4	US-09-617-053A-4	Sequence 4, Appli
31	909	27.7	497	4	US-09-201-936-4	Sequence 4, Appli
32	908	27.7	497	4	US-08-657-755-2	Sequence 2, Appli
33	874	26.7	496	2	US-08-511-485-10	Sequence 10, Appl
34	874	26.7	496	3	US-09-212-971-10	Sequence 10, Appl
35	874	26.7	496	3	US-08-800-929A-10	Sequence 10, Appl
36	874	26.7	496	4	US-09-617-053A-10	Sequence 10, Appl
37	874	26.7	497	4	US-09-201-936-10	Sequence 10, Appl
38	737	22.5	498	4	US-08-657-755-1	Sequence 1, Appl
39	737	22.5	498	2	US-09-493-784-22	Sequence 22, Appl
40	736.5	22.5	498	3	US-08-836-134-22	Sequence 22, Appl
41	735.5	22.5	497	3	US-08-511-485-13	Sequence 13, Appl
42	513	15.7	268	4	US-09-201-936-13	Sequence 13, Appl
43	513	15.7	268	4	US-09-239-867-4	Sequence 40, Appl
44	492.5	15.0	236	4	US-09-502-528-3	Sequence 4, Appl
45	466.5	14.2	377	4	US-08-836-134-23	Sequence 3, Appl
	463	14.1	1151	3		Sequence 23, Appl

ALIGNMENTS

RESULT 1
US-08-569-749-2
; Sequence 2, Application US/0856974 9
; Patent No. 6187557
; GENERAL INFORMATION:
; APPLICANT: Rothe, Mike
; APPLICANT: Goedel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC Compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749
; APPLICATION NUMBER:
; FILING DATE:
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Brezner, David J.
; REGISTRATION NUMBER: 24,774
; REFERENCE/DOCKET NUMBER: A-62464/DJB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415)781-1989
; TELEFAX: (415)398-3249
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 618 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-569-749-2

Query Match   100.0%; Score 3277; DB 3; Length 618;
Best Local Similarity   100.0%; Pred. No. 4.8e-308;
Matches 618; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKTDFSCELYRMSTYSTFPAGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

RESULT 2

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US-09-069-023-29
Sequence 29, Application US/0969023A
Patent No. 6348573
GENERAL INFORMATION:
Patent No. 6348573
APPLICANT: Nunez, Gabriel
APPLICANT: Inohara, Naohiro
APPLICANT: Koseki, Takeyoshi
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR IDENTIFYING APOPTOSIS
TITLE OF INVENTION: SIGNALING PATHWAY INHIBITORS AND ACTIVATORS
FILE REFERENCE: UM-03333
CURRENT APPLICATION NUMBER: US/09/069,023A
CURRENT FILING DATE: 1998-04-27
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 29
LENGTH: 618
TYPE: PRT
ORGANISM: Homo sapiens
US-09-069-023-29

Query Match      100.0%;    Score 3277;    DB 4;    Length 618;
Best Local Similarity    100.0%;    Pred. No. 4.8e-308;
Matches  618;  Conservative  0;  Mismatches  0;  Indels  0;  Gaps  0;
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Qy   61 PVSERSLARAGFYYTGVNDKVXCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
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Db   61 PVSERSLARAGFYYTGVNDKVXCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
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Qy  181 MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
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Db  181 MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
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Db  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
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Db  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy  361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
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Db  361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420

Qy  421 GENYKTVNDIVSALLNAEDEKREBEEKQAEBMASDDLSLIRKNRMALFQQLTCVLPILD 480
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Db  601 KCPICRGIIKGTVRTFLS 618
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RESULT 3

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PCT-US96-12860-2
Sequence 2, Application PCT/US9612860
GENERAL INFORMATION:
APPLICANT: TULARIK, INC.
TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/12860
FILING DATE: 06 AUG 1996
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U.S. Serial Nos. 08/512,946 & 08/569,749
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Brezner, David J.
REGISTRATION NUMBER: 24,774
REFERENCE/DOCKET NUMBER: A-62464/DJB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415)781-1989
TELEFAX: (415)398-3249
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 618 amino acids
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Qy  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
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Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
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Db  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
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Qy  361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
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Qy  421 GENYKTVNDIVSALLNAEDEKREBEEKQAEBMASDDLSLIRKNRMALFQQLTCVLPILD 480
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Db  421 GENYKTVNDIVSALLNAEDEKREBEEKQAEBMASDDLSLIRKNRMALFQQLTCVLPILD 480

Qy  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKBIDSTLYKNLF 540
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKBIDSTLYKNLF 540

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Db  541 VDKNMKYIPTEDVSGLSLEEQLRRLQEBRTCKVCMDKEVsVVFIPCGHLVVCQEECAPSLR 600

Qy  601 KCPICRGIIKGTVRTFLS 618
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Db  601 KCPICRGIIKGTVRTFLS 618
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; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
PCT-US96-12860-2

Query Match   100.0%; Score 3277; DB 5; Length 618;
Best Local Similarity 100.0%; Pred. No. 4.8e-308;
Matches 618; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMIQDFSCELYRMSTYSTFPPAGV 60
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Db   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMIQDFSCELYRMSTYSTFPPAGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy   121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNFLNSRAVEDISSSRTNPYSYA 180
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Db   121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNFLNSRAVEDISSSRTNPYSYA 180

Qy   181 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACCGGKLSNWEPKDDAMSEH 240
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Db   181 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACCGGKLSNWEPKDDAMSEH 240

Qy   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy   301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
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Db   301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy   361 STSDTTGEENADPPIIHFGPGEBSSSEDAVMMNTPVVKSALIMGFNRDLVKQTVQSKILTT 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   361 STSDTTGEENADPPIIHFGPGEBSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420

Qy   421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLLSLIRKNRMALFQQLTCVLPILD 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLLSLIRKNRMALFQQLTCVLPILD 480

Qy   481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 540
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Db   481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 540

Qy   541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
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Db   541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy   601 KCPICRGIIKGTVRTFLS 618
          ||||||||||||||||||
Db   601 KCPICRGIIKGTVRTFLS 618

RESULT 4
US-08-511-485-8
; Sequence 8, Application US/08511485
; Patent No. 5919912
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mackenzie, Alexander E.
; APPLICANT: Baird, Stephen
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; NUMBER OF SEQUENCES: 38
; CORRESPONDENCE ADDRESS:
;   ADDRESSEE: Fish & Richardson P.C.
;   STREET: 225 Franklin Street
;   CITY: Boston
;   STATE: MA
;   COUNTRY: USA
;   ZIP: 02110-2804

; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/511,485
; FILING DATE: 04-AUG-1995
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Clark, Paul T.
; REGISTRATION NUMBER: 30,162
; REFERENCE/DOCKET NUMBER: 07540/002001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617/542-5070
; TELEFAX: 617/542-8906
; TELEX: 200154
; INFORMATION FOR SEQ ID NO: 8:
; SEQUENCE CHARACTERISTICS:
;   LENGTH: 618 amino acids
;   TYPE: amino acid
;   STRANDEDNESS: not relevant
;   TOPOLOGY: both
; MOLECULE TYPE: protein
US-08-511-485-8

Query Match   99.1%; Score 3247; DB 2; Length 618;
Best Local Similarity 99.4%; Pred. No. 3.8e-305;
Matches 614; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

Qy   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMIQDFSCELYRMSTYSTFPPAGV 60
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Db   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMIQDFSCELYRMSTYSTFPPAGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
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Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy   121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPPNFLNSRAVEDISSSRTNPYSYA 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPPNFLNSRAVEDISSSRTNPYSYA 180

Qy   181 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACCGGKLSNWEPKDDAMSEH 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
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Db   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy   301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
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Db   301 NDDVKCFGCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy   361 STSDTTGEENADPPIIHFGPGEBSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
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Db   361 STSDTTGEENADPPIIHFGPGEBSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVLSKILTT 420

Qy   421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLLSLIRKNRMALFQQLTCVLPILD 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLLSLIRKNRMALFQQLTCVLPILD 480

Qy   481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 540
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIWVKGNAAANIFKNCLKEIDSTLYKNLF 540

Qy   541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
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Db   541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy   601 KCPICRGIIKGTVRTFLS 618
          ||||||||||||||||||
Db   601 KCPICRGIIKGTVRTFLS 618

RESULT 5
US-09-212-971-8
; Sequence 8, Application US/09212971B
; Patent No. 6107041
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009002
; CURRENT APPLICATION NUMBER: US/09/212,971B
; CURRENT FILING DATE: 1998-12-16
; EARLIER APPLICATION NUMBER: 60/017,354
; EARLIER FILING DATE: 1996-04-26
; EARLIER APPLICATION NUMBER: 60/030,590
; EARLIER FILING DATE: 1996-11-14
; EARLIER APPLICATION NUMBER: 08/800,929
; EARLIER FILING DATE: 1997-02-13
; NUMBER OF SEQUENCES: 17
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-212-971-8

Query Match   99.1%;   Score 3247; DB 3;   Length 618;
Best Local Similarity 99.4%; Pred. No. 3.8e-305;
Matches 614; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy    1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV  60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV  60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180

Qy  181 MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVAC?ACGGKLSNWEPKDDAMSEH 240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  181 MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVAC?ACGGKLSNWBPKDDAMSEH 240

Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAAARMRTFMYWPSSVQPBQLASAGFYYVGR 300
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAAARMRTFMYWPSSVQPBQLASAGFYYVGR 300

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLBQLL 360
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLBQLL 360

Qy  361 STSDTTGBENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
        |||||||||||||||||||||||||||||||||||||||||||||||||||| |||||||
Db  361 STSDTTGBENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVLSKILTT 420

Qy  421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD 480
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD 480

Qy  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTLLVKGNAAANIFKNCLKBIDSTLYKNLF 540
        ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIWVKGNAAANIFKNCLKBIDSTLYKNLF 540

Qy  541 VDKNMKYIPTEDVSGLSLBEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  541 VDKNMKYIPTEDVSGLSLBEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy  601 KCPICRGIIKGTVRTFLS 618
        ||||||||||||||||||
Db  601 KCPICRGIIKGTVRTFLS 618
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RESULT 6
US-08-800-929A-8
; Sequence 8, Application US/08800929A
; Patent No. 6133437
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF
; TITLE OF INVENTION: IAPS AND NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIV
; TITLE OF INVENTION: DISEASE
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Clark & Elbing LLP
; STREET: 176 Federal Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/800,929A
; FILING DATE: 13-FEB-1997
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/030,590
; FILING DATE: 14-NOV-1996
; APPLICATION NUMBER: 60/017,354
; FILING DATE: 26-APR-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Bieker-Brady, Kristina
; REGISTRATION NUMBER:
; REFERENCE/DOCKET NUMBER: 07891/009001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-428-0200
; TELEFAX: 617-428-7045
; TELEX:
; INFORMATION FOR SEQ ID NO: 8:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 618 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-800-929A-8

Query Match   99.1%;   Score 3247; DB 3;   Length 618;
Best Local Similarity 99.4%; Pred. No. 3.8e-305;
Matches 614; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy    1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV  60
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Db    1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV  60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
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Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy  541 VDKNMKYIPTEDVSGLSLBEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  541 VDKNMKYIPTEDVSGLSLBEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy  601 KCPICRGIIKGTVRTFLS 618
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Db  601 KCPICRGIIKGTVRTFLS 618
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RESULT 7
US-09-617-053A-8
; Sequence 8, Application US/09617053A
; Patent No. 6300492
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009003
; CURRENT APPLICATION NUMBER: US/09/617,053A
; CURRENT FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 08/800,929
; PRIOR FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SEQ ID NO 8
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
; SOFTWARE: FastSEQ for Windows Version 4.0
US-09-617-053A-8

Query Match   99.1%; Score 3247; DB 4; Length 618;
Best Local Similarity   99.4%; Pred. No. 3.8e-305;
Matches 614; Conservative   0; Mismatches   4; Indels   0; Gaps   0;

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Qy    1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQXMKYDFSCELYRMSTYSTPPAGV 60
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Db    1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQXMKYDFSCELYRMSTYSTPPAGV 60

Qy  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
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Db  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180

Qy  181 MSTEEARFLITYHMWPLITFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  181 MSTEEARFLITYHMWPLITFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWFSSVPVQPEQLASAGFYYVGR 300
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWFSSVPVQPEQLASAGFYYVGR 300

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy  361 STSDTTGEENADPPIIHFGPGBSSSSEDAVMMNTPVVRSALEMGFNRDLVKQTVQSKILTT 420
        |||||||||||||||||||||||||||||||||||||||||||||||||||| ||||||
Db  361 STSDTTGEENADPPIIHFGPGBSSSSEDAVMMNTPVVRSALEMGFNRDLVKQTVLSKILTT 420

Qy  421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLLSLIRKNRMALFQQLTCVLPILD 480
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLLSLIRKNRMALFQQLTCVLPILD 480

Qy  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKBIDSTLYKNLF 540
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Db  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIWVKGNAAANIFKNCLKBIDSTLYKNLF 540

Qy  541 VDKNMKYIPTEDVSGLSLEEQLRRLQBERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  541 VDKNMKYIPTEDVSGLSLEEQLRRLQBERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy  601 KCPICRGIIKGTVRTFLS 618
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Db  601 KCPICRGIIKGTVRTFLS 618
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RESULT 8
US-09-201-936-8
; Sequence 8, Application US/09201936
; Patent No. 6541457
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: MacKenzie, Alexander E.
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; FILE REFERENCE: 07891/003003
; CURRENT APPLICATION NUMBER: US/09/201,936
; CURRENT FILING DATE: 1998-12-01
; EARLIER APPLICATION NUMBER: 09/011,356
; EARLIER FILING DATE: 1998-02-04
; EARLIER APPLICATION NUMBER: PCT/IB96/01022
; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 08/576,956
; EARLIER FILING DATE: 1995-12-22
; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SEQ ID NO 8
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
; SOFTWARE: FastSEQ for Windows Version 3.0
US-09-201-936-8

Query Match   99.1%; Score 3247; DB 4; Length 618;
Best Local Similarity   99.4%; Pred. No. 3.8e-305;

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Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
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Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180

Qy  181 MSTEEARFLITYHMWPLITFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  181 MSTEEARFLITYHMWPLITFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWFSSVPVQPEQLASAGFYYVGR 300
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWFSSVPVQPEQLASAGFYYVGR 300

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy  361 STSDTTGEENADPPIIHFGPGBSSSSEDAVMMNTPVVRSALEMGFNRDLVKQTVQSKILTT 420
        |||||||||||||||||||||||||||||||||||||||||||||||||||| ||||||
Db  361 STSDTTGEENADPPIIHFGPGBSSSSEDAVMMNTPVVRSALEMGFNRDLVKQTVLSKILTT 420

Qy  421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLLSLIRKNRMALFQQLTCVLPILD 480
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLLSLIRKNRMALFQQLTCVLPILD 480

Qy  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKBIDSTLYKNLF 540
        |||||||||||||||||||||||||||||||| |||||||||||||||||||||||||||
Db  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIWVKGNAAANIFKNCLKBIDSTLYKNLF 540

Qy  541 VDKNMKYIPTEDVSGLSLEEQLRRLQBERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
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Db  541 VDKNMKYIPTEDVSGLSLEEQLRRLQBERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy  601 KCPICRGIIKGTVRTFLS 618
        ||||||||||||||||||
Db  601 KCPICRGIIKGTVRTFLS 618
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RESULT 9
US-08-569-749-14
; Sequence 14, Application US/0869749
; Patent No. 6187557
; GENERAL INFORMATION:
; APPLICANT: Rothe, Mike
; APPLICANT: Goeddel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749
; FILING DATE:
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:

Matches 614; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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Qy    1 MKHTASQRLFPGPSYQNIKSIMEDSTILLSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV   60
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1 MKHTASQRLFPGPSYQNIKSIMEDSTILLSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV   60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS  120
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Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS  120

Qy  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA  180
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Db  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA  180

Qy  181 MSTEEARFLITYHNWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH  240
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Db  181 MSTEEARFLITYHNWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH  240

Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMTWPSSVPVQPEQLASAGFYYVGR  300
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMTWPSSVPVQPEQLASAGFYYVGR  300

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL  360
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL  360

Qy  361 STSDTTGEENADPPIIHFGPGEBSSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILT  420
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  361 STSDTTGEENADPPIIHFGPGEBSSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILT  420

Qy  421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSIIRKNRMALFQQLTCVLPILD  480
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSIIRKNRMALFQQLTCVLPILD  480

Qy  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF  540
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF  540

Qy  541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR  600
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR  600

Qy  601 KCPICRGIIKGTVRTFLS  618
        ||||||||||||||||||
Db  601 KCPICRGIIKGTVRTFLS  618
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NAME: Brezner, David J.
REGISTRATION NUMBER: 24,774
REFERENCE/DOCKET NUMBER: A-62464/DJB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415)781-1989
TELEFAX: (415)398-3249
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 612 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-569-749-14

Query Match 83.2%; Score 2728; DB 3; Length 612;
Best Local Similarity 83.4%; Pred. No. 6.2e-255;
Matches 517; Conservative 45; Mismatches 48; Indels 10; Gaps 6;

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Qy    1 MKHTASQRLFPGPSYQNIKSIMEDSTILLSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV   60
        |||||||||||:|||  |:| ||||||||||||||||||||||||||||||||||||||||
Db    1 MDKTVSQRLQGQGTLHQKLKRIMEKSTILSNWTKESEBKKMKPDFSCELYRMSTYSAFPRGV   60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS  120
        ||||||||||||||||:|||| ::|||||||||||||||||: ::|||| :||||| |||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPVEKHRQFYPSCSFVQTLLSAS  120

Qy  121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA  180
        ||||||||||||||: |||||||-||---------||||||||||||||:|||||||||||
Db  121 LQSPSKNMSPVKSRFAHS-SP-LER----GGIHSNLCSSPLNSRAVEDF-SSRMDPCSYA  173

Qy  181 MSTEEARFLITYHNWPLTFLSPSPAELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH  240
        ||||||||||:||||||||||:| |||:||||||||||||||||||||||||||||||||
Db  174 MSTEEARFLTYSMWPLSFSLSPAELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH  233

Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMTWPSSVPVQPEQLASAGFYYVGR  300
        ||||:|||||||::| ::|:|||||||||::|||||||:|||||||||||:|||||||||
Db  234 RRHFPHCPFLENTSETQRFSISNLSMQTSARLRTFLYWPPSVPVQPEQLASAGFYYVGR  293

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL  360
        ||||||||||||||||| ||||::|||||||||||||||||||||||||||:|||||||
Db  294 NDDVKCFCCDGGLRCWEPGDDPWIEHAKWFPRCEFLIRMKGQEFVDEIQARYPHLLEQLL  353

Qy  361 STSDTTGEENADP--PIIHFGPGEBSSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKIL  418
        ||||||||||||   ||||:||||| ::||||||||||||||||||||||||:|||||||:
Db  354 STSDTPGEENADPTETVVHFGPGE-SSEDVVMMSTPVVKAALEMGFSRSLVRQTVQRQIL  412
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RESULT 10
PCT-US96-12860-14
; Sequence 14, Application PC/TUS9612860
; GENERAL INFORMATION:
; APPLICANT: TULARIK, INC.
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT

```
Qy  419 TTGENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPI  478
        ||||||||||||||||||||||||:||:||||| ::||||||:|||||||||||||||||
Db  413 ATGENYRTVNDIVSVLLNAEDERREEEKERQTBENASGDLSIIRKNRMALFQQLTHVLPI  472

Qy  479 LDNLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKN  538
        ||:||:||:|||||||||:||||||||||||||||||||||||||:||:|||||||||||
Db  473 LDNLLRASVITKQEHDIIRQKTQIPLQARELIDTVLVKGNAAANIFNSLKEIDSTLYEN  532

Qy  539 LFVDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPS  598
        ||:||||||||||||||||||||||||||||||||||||:|||||||||||||||||||
Db  533 LFVEKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDREVSIVFIPCGHLVVCQECAPS  592

Qy  599 LRKCPICRGIIKGTVRTFLS  618
        ||||||||||:|||||||||
Db  593 LRKCPICRGTIKGTVRTFLS  612
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```
         STREET:  4 Embarcadero Center, Suite 3400
         CITY:  San Francisco
         STATE:  California
         COUNTRY:  USA
         ZIP:  94111
     COMPUTER READABLE FORM:
         MEDIUM TYPE:  Floppy disk
         COMPUTER:  IBM PC compatible
         OPERATING SYSTEM:  PC-DOS/MS-DOS
         SOFTWARE:  Patentin Release #1.0, Version #1.30)
     CURRENT APPLICATION DATA:
         APPLICATION NUMBER:  PCT/US96/12860
         FILING DATE:  06 AUG 1996
         CLASSIFICATION:
     PRIOR APPLICATION DATA:
         APPLICATION NUMBER:  U.S. Serial Nos. 08/512,946 & 08/569,749
         CLASSIFICATION:
     ATTORNEY/AGENT INFORMATION:
         NAME:  Brenner, David J.
         REGISTRATION NUMBER:  24,774
         REFERENCE/DOCKET NUMBER:  A-62464/DJB
     TELECOMMUNICATION INFORMATION:
         TELEPHONE:  (415)781-1989
         TELEFAX:  (415)398-3249
 INFORMATION FOR SEQ ID NO: 14:
     SEQUENCE CHARACTERISTICS:
         LENGTH:  612 amino acids
         TYPE:  amino acid
         STRANDEDNESS:  single
         TOPOLOGY:  linear
         MOLECULE TYPE:  protein
PCT-US96-12860-14

Query Match            83.2%;  Score 2728;  DB 5;  Length 612;
Best Local Similarity  83.4%;  Pred. No. 6.2e-255;
     Matches 517;  Conservative 45;  Mismatches 48;  Indels 10;  Gaps 6;

Qy    1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60
          |||  ||| :|  :| | ||||:|| |||||||||||||||||||||||||||||
Db    1 MDKTVSQRLGQGTLHQKLKRIMEKSTILSNWTKESEEKMKPDFSCELYRMSTYSAPPRGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
         ||||||||||||||||||||||||||||||   :|||||:|||||||||||||||||||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKQGDSPVEKHRQFYPSCSFVQTLLSAS 120

Qy  121 LGSTSKQNTSPMRNSPAHSLSPTLEHSSLFSGSYSSLSSPNPLNSRAVEDISSSRTNPYSYA 180
         |||||| : :| ||||||||||||||||||||||||||||||||||||||| | ||||||
Db  121 LQSPSKNMSPVKSRFAHS-SP-LER----GGIHSNLCSSPLNSRAVEDF-SSRMDPCSYA 173

Qy  181 MSTEEARFLITYHMWPLITFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  174 MSTEEARFLITYSMWPLSFLSPAELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 233

Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
         |||||||||||| : :||||||||||||||||:||||||||||||||||||||||||||
Db  234 RRHFPHCPFLENTSETQRFSISNLSMQTHSARLRTFLYWPPSVPVQPEQLASAGFYYVDR 293

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  294 NDDVKCFCCDGGLRCWEPGDDPWIEHAKWFPRCEFLIRMKGQ3FVDEIQARYPHLLEQLL 353

Qy  361 STSDTTGEENADP--PIIHGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKIL 418
         |||||:||||||   : :|||||||||||||||| ||||||||||||||||::|||||
Db  354 STSDTPGEENADPTETVVHFGPGE-SSBDVVMMSTPVVKAALEMGFSRSLVRQTVQROIL 412

Qy  419 TTGENYKIVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSL:RKNRMALFQOLTCVLPI 478
         |||||||||||||||||||||||||:||||||||||||||||:|||||||||||||||
Db  413 ATGENYRTVNDIVSVLLNAEDEREEEKERQTEEMASGDLSLIRKNRMALFQQLTHVLPI 472

Qy  479 LDNLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKN 538
         |||||||:||| |||| : : ||||||||||||||||||||||  |||| ||||:|| :|
Db  473 LDNLLEASVTKQEHDIIIROKTQIPLQARELIDTVLVKGNAAANIFKNSLKEIDSTLYEN 532
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```
RESULT 11
US-09-212-971-14
 Sequence 14, Application US/09212971B
 Patent No. 6107041
 GENERAL INFORMATION:
     APPLICANT:  Korneluk, Robert G
     APPLICANT:  MacKenzie, Alexander E
     APPLICANT:  Liston, Peter
     APPLICANT:  Baird, Stephen
     APPLICANT:  Tsang, Benjamin K
     APPLICANT:  Pratt, Christine
     TITLE OF INVENTION:  DETECTION AND MODULATION OF IAPS AND
     TITLE OF INVENTION:  NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
     TITLE OF INVENTION:  DISEASE
     FILE REFERENCE:  07891/009002
     CURRENT APPLICATION NUMBER:  US/09/212,971B
     CURRENT FILING DATE:  1998-12-16
     EARLIER APPLICATION NUMBER:  60/017,354
     EARLIER FILING DATE:  1996-04-26
     EARLIER APPLICATION NUMBER:  60/030,590
     EARLIER FILING DATE:  1996-11-14
     EARLIER APPLICATION NUMBER:  08/800,929
     EARLIER FILING DATE:  1997-02-13
     NUMBER OF SEQ ID NOS:  17
     SOFTWARE:  FastSEQ for Windows Version 4.0
 SEQ ID NO 14
     LENGTH:  612
     TYPE:  PRT
     ORGANISM:  Mus musculus
US-09-212-971-14

Query Match            83.1%;  Score 2724;  DB 3;  Length 612;
Best Local Similarity  83.2%;  Pred. No. 1.5e-254;
     Matches 516;  Conservative 46;  Mismatches 48;  Indels 10;  Gaps 6;

Qy    1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60
          |||  ||| :|  :| | ||||:|| |||||||||||||||||||||||||||||
Db    1 MDKTVSQRLGQGTLHQKLKRIMEKSTILSNWTKESEEKMKPDFSCELYRMSTYSAPPRGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
         ||||||||||||||||||||||||||||||   :|||||:|||||||||||||||||||
Db    1 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKQGDSPVEKHRQFYPSCSFVQTLLSAS 120

Qy  121 LGSTSKQNTSPMRNSPAHSLSPTLEHSSLFSGSYSSLSSPNPLNSRAVEDISSSRTNPYSYA 180
         |||||| : :| ||||||||||||||||||||||||||||||||||||||| | ||||||
Db  121 LQSPSKNMSPVKSRFAHS-SP-LER----GGIHSNLCSSPLNSRAVEDF-SSRMDPCSYA 173

Qy  181 MSTEEARFLITYHMWPLITFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  174 MSTEEARFLITYSMWPLSFLSPAELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy  241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
         |||||||||||| : :||||||||||||||||:||||||||||||||||||||||||||
Db  234 RRHFPHCPFLENTSETQRFSISNLSMQTHSARLRTFLYWPPSVPVQPEQLASAGFYYVDR 293

Qy  301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  294 NDDVKCFCCDGGLRCWEPGDDPWIEHAKWFPRCEFLIRMKGQ3FVDEIQARYPHLLEQLL 360

Qy  361 STSDTTGEENADP--PIIHGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKIL 418
         |||||:||||||   : :|||||||||||||||| ||||||||||||||||::|||||
Db  354 STSDTPGEENADPTETVVHFGPGE-SSKDVVMMSTPVVKAALEMGFSRSLVRQTVQROIL 412

Qy  419 TTGENYKIVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPI 478
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RESULT 12
US-08-800-929A-14
Sequence 14, Application US/08800929A
Patent No. 6133437
GENERAL INFORMATION:
    APPLICANT: Korneluk, Robert G
    APPLICANT: MacKenzie, Alexander E
    APPLICANT: Liston, Peter
    APPLICANT: Baird, Stephen
    APPLICANT: Tsang, Benjamin K
    APPLICANT: Pratt, Christine
    TITLE OF INVENTION: DETECTION AND MODULATION OF
    TITLE OF INVENTION: IAPS AND NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
    TITLE OF INVENTION: DISEASE
    NUMBER OF SEQUENCES: 17
    CORRESPONDENCE ADDRESS:
        ADDRESSEE: Clark & Elbing LLP
        STREET: 176 Federal Street
        CITY: Boston
        STATE: MA
        COUNTRY: USA
        ZIP: 02110
    COMPUTER READABLE FORM:
        MEDIUM TYPE: Diskette
        COMPUTER: IBM Compatible
        OPERATING SYSTEM: DOS
        SOFTWARE: FastSEQ for Windows Version 2.0
    CURRENT APPLICATION DATA:
        APPLICATION NUMBER: US/08/800,929A
        FILING DATE: 13-FEB-1997
        CLASSIFICATION: 424
    PRIOR APPLICATION DATA:
        APPLICATION NUMBER: 60/030,590
        FILING DATE: 14-NOV-1996
        APPLICATION NUMBER: 60/017,354
        FILING DATE: 26-APR-1996
    ATTORNEY/AGENT INFORMATION:
        NAME: Bieker-Brady, Kristina
        REGISTRATION NUMBER:
        REFERENCE/DOCKET NUMBER: 07891/009001
    TELECOMMUNICATION INFORMATION:
        TELEPHONE: 617-428-0200
        TELEFAX: 617-428-7045
        TELEX:
INFORMATION FOR SEQ ID NO: 14:
    SEQUENCE CHARACTERISTICS:
        LENGTH: 612 amino acids
        TYPE: amino acid
        STRANDEDNESS: single
        TOPOLOGY: linear
        MOLECULE TYPE: protein
US-08-800-929A-14

Query Match   83.1%;   Score 2724; DB 3; Length 612;
Best Local Similarity   83.2%;   Pred. No. 1.5e-254;
Matches 516; Conservative   46; Mismatches 48; Indels 10; Gaps   6;

Qy   413 ATGENYRTVNDIVSVLLNAEDERREEEKERQTEEMASGDLSLIRKNRMALFQQLTHVLPI 472
           ||||||||||||||||||||||||||||||||||||||||||||||||||||| ||||
Db   413 ATGENYRTVNDIVSVLLNAEDERREEEKERQTEEMASGDLSLIRKNRMALFQQLTHVLPI 472

Qy   473 LDNLLEASVITKQEHDIIROKTQIPLQARELIDTVLVKGNAAAANIFKNSLKEIDSTLYEN 532
           |||||||||||||||||||||||:||||||||||||||||||||||||||||||||||||
Db   473 LDNLLEASVITKQEHDIIROKTQIPLQARELIDTVLVKGNAAAANIFKNSLKEIDSTLYEN 532

Qy   533 LFVEKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDIEVSVVFIPCGHLVVCQECAPS 592
           |||||||||||||||||||||||||||||||||||: ||||||||||||||||||||||
Db   533 LFVEKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMD:EVSIVFIPCGHLVVCQECAPS 592

Qy   593 LRKCPICRGIIKGTVRTFLS 612
           ||||||||||||||||||||
Db   593 LRKCPICRGIIKGTVRTFLS 612

Qy   599 LRKCPICRGTIKGTVRTFLS 618
           ||||||||::||||||||||

RESULT 13
US-09-617-053A-14
Sequence 14, Application US/0961705JA
Patent No. 6300492
GENERAL INFORMATION:
    APPLICANT: Korneluk, Robert G
    APPLICANT: MacKenzie, Alexander E
    APPLICANT: Liston, Peter
    APPLICANT: Baird, Stephen
    APPLICANT: Tsang, Benjamin K
    APPLICANT: Pratt, Christine
    TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
    TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
    TITLE OF INVENTION: DISEASE
    FILE REFERENCE: 07891/009003
    CURRENT APPLICATION NUMBER: US/09/617,053A
    CURRENT FILING DATE: 2000-07-14
    PRIOR APPLICATION NUMBER: US 08/800,929
    PRIOR FILING DATE: 1997-02-13
    NUMBER OF SEQ ID NOS: 17
    SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14
    LENGTH: 612
    TYPE: PRT
    ORGANISM: Mus musculus
US-09-617-053A-14

Query Match   83.1%;   Score 2724; DB 4; Length 612;

Qy   1 MHKTASQRLFPGPSYQNIKSIMEDSTILLSDWTNSNKQKMKYDPSCELYRMSTYSTFPAGY 60
           |:|||:||||||||||||||||||||:|||||||||||||||:||||||||||||:||||
Db   1 MDKTVSQRLQGQTLHQLKRIMEKSTILLSNWTKSSEBKMKFDPSCELYRMSTYSAFPRGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
           ||||||||||||||||||||||||||||||||||||||||:|||:|||||||||:|||||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKGDSPVEKHRQFYPSCSFVQTLLSAS 120

Qy   121 LGSTSKNTSPMRNSPAHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
           |||||||:||||||||||||:||||||||||:||::|:|||||||||||||||:||||||
Db   121 LQSPSKNMSPVKSRFAHS-SP-LER----GGIHSNLCSSPLNSRAVEDF-SSRMDPCSYA 173

Qy   181 MSTEBARPLTYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWBPKDDAMSEH 240
           ||||||||| |||||||:|||||||||||||||||||||||||||||||||||:|||||
Db   174 MSTEBARFLTYSMWPLSFLSPAELARAGFYYIGPGDRVACFACGGKLSNWBPKDDAMSEH 233

Qy   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAAARVRTFMYWPSSVPVQPEQLASAGFYYVGR 300
           |||||||||||:||||||||||||:|||||:||||||||||||:||:|||||||||||||
Db   234 RRHFPHCPFLENTSETQRFSISNLSMQTHSARLRTFLYWPPSVPVQPEQLASAGFYYVDR 293

Qy   301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQBFVDEIQGRYPHLLEQLL 360
           |||||||||||||||||||||||||||||||||||||||||||:|||||:||||||||||
Db   294 NDDVKCFCCDGGLRCWEPGDDPWIEHAKWFPRCEFLIRMKGQBFVDEIQARYPHLLEQLL 353

Qy   361 STSDTTGBENADP--PIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKIL 418
           |||||||||||| :||||||||::||||||||||||||:|||||||||||:|||||:|||
Db   354 STSDTPGBENADPTETVVHFGPGE-SSKDVVMMSTPVVKAALEMGFSRSLVRQTVQROIL 412

Qy   419 TTGENYKTVNDIVSALLNAEDEKRBEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPI 478
           |||||||||||||:|||||||||:|||||||:||||||||||||||||||||||||||||
Db   413 ATGENYRTVNDIVSVLLNAEDERREEEKERQTEEMASGDLSLIRKNRMALFQQLTHVLPI 472

Qy   479 LDNLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAAANIFKNCLKEIDSTLYKN 538
           |||||:|||||||||||:||||||||||||||:|||||||||||||||||:|||||||||
Db   473 LDNLLEASVITKQEHDIIROKTQIPLQARELIDTVLVKGNAAAANIFKNSLKEIDSTLYEN 532

Qy   539 LFVDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPS 598
           |||:|||||||||||||||||||||||||||||||||||:||||||||||||||||||||
Db   533 LFVEKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDREVSIVFIPCGHLVVCQECAPS 592

Qy   599 LRKCPICRGIIKGTVRTFLS 618
           |||||||||||||||||||
Db   593 LRKCPICRGTIKGTVRTFLS 612

RESULT 14
US-09-201-936-42
; Sequence 42, Application US/0920193б
; Patent No. 6541457
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: MacKenzie, Alexander E.
; APPLICANT: Baird, Stephen
; APPLICANT: Liston, Peter
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; FILE REFERENCE: 07891/003003
; CURRENT APPLICATION NUMBER: US/09/201,936
; CURRENT FILING DATE: 1998-12-01
; EARLIER APPLICATION NUMBER: 09/011,356
; EARLIER FILING DATE: 1998-02-04
; EARLIER APPLICATION NUMBER: PCT/IB96/01022
; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 08/576,956
; EARLIER FILING DATE: 1995-12-22
; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 42

Best Local Similarity 83.2%; Pred. No. 1.5e-254;
Matches 516; Conservative 46; Mismatches 48; Indels 10; Gaps 6;

Qy   1 MEKTASQRLFPGPSYSYQNIKSIMEDSTILLSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1 MDKTVSQRLGQGTLHQKLKRIMEKSTILLSNWTLKESEERKMKPDFSCELYRMSTYSAFPRGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFPCCGGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
           |||||:|||||||||||||||||||||:||||||||||:||:||:|:|||||||||:|:|||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGGLMLDNWKQGDSPVEKHRQFYPSCSFVQTLLSAS 120

Qy   121 LQSPSKNMSPVKSRFAHS-SP-LER-----GGIHSNLCSSPLNSRAVEDF-SSRMDPCSYA 173
           |:|||||||||||||||| :|: ||| ||||||||:|||||||||| ::||:||||
Db   121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLPSGSYSSLSPNPLNSRAVEDISSSRTNPSYSYA 180

Qy   174 MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVAC:FACGGGKLSNWEPKDDAMSEH 233
           |||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||||
Db   181 MSTEEARFLTYSMWPLSFLSPAELARAGFYYIGPGDRVAC:FACGGGKLSNWEPKDDAMSEH 240

Qy   234 RRHFPHCPFLENTSETQRFSISNLSMQTHSARLRTFLYWPPSVPVQPEQLASAGFYYVDR 293
           |||||||||||||||||||:||||||||||||||||:||||||||||||||||||||||||
Db   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWISSVPVQPEQLASAGFYYVGR 300

Qy   294 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 353
           |||||||||||||||||||||||||||||||||||||||:||||||:|||:|:||||||||
Db   301 NDDVKCFCCDGGLRCWESGDDPWIEHAKWFPRCEFLIRMKGQEFVDEIQARYPHLLEQLL 360

Qy   354 STSDTPGEENADPTETVVHFGPGE--SSKDVVMMSTPVVKALEMGFSRSLVRQTVQROIL 412
           :|||||||||||||||||||||| ||:|||||||||||||||:||||||||||||:||
Db   361 STSDTTGEENADP--PIIHFGPGESSSSEDAVMMNTPVVKSALEMGPNRDLVKQTVQSKIL 418

Qy   413 ATGENYRTVANDIVSVLLNAEDERREEEKEKQAAEMASDLLSLIRKNRMALFQQLTCVLPI 472
           :|||||||||||||||||||||||||||||||||||||||:|||||||||||||||||||
Db   419 TTGENYKTVANDIVSALLNAEDERKREEEKEKQAEMASDLLSLIRKNRMALFQQLTVLPI 478

Qy   473 LDNLLEASVITKQEHDIIRQKTOIPLQARELIDTVLVKGNVNAANIFKNSLKEIDSTLYEN 532
           |||||||:||||||||||:||: |||||||||:||||||||:|||||||:|||:|||||||
Db   479 LDNLLKANVINKQEHDIIKQKTOIPLQARELIDTILVKGNXNAANIFKNCLKEIDSTLYKN 538

Qy   533 LFVEKNMKYIPTEDVSGLSLEEQLRRLQEBRTCKVCMDKEVSVVFIPCGHLVVCQEBCAPS 592
           ||:|||||||||||||||||||||||||||||||||||||:|||:|||||||||||||||
Db   539 LFVDKNMKYIPTEDVSGLSLREQLRRLQEBRTCKVCMDRVSIVFIPCGHLVVCQEBCAPS 598

Qy   593 LRKCPICRGIIKGTVRTFLS 612
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Db   599 LRKCPICRGIIKGTVRTFLS 618

;
; LENGTH: 591
; TYPE: PRT
; ORGANISM: Mus musculus
US-09-201-936-42

RESULT 15
US-08-569-749-4
; Sequence 4, Application US/08569749
; Patent No. 6187557
; GENERAL INFORMATION:
; APPLICANT: Rothe, Mike
; APPLICANT: Goeddel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749

Query Match   83.8%;
Best Local Similarity 83.8%; Score 2654; DB 4; Length 591;
Matches 502; Conservative 44; Mismatches 43; Indels 10; Gaps 6;

Qy   22 MEDSTILLSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGVPVSERSLARAGFYYTGVNDKV 81
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Db   1 MEKSTILLSNWTLKESEERKMKPDFSCELYRMSTYSAFPRGVPVSERSLARAGFYYTGVNDKV 60

Qy   82 KCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSASLGSTSKNTSPMRNSFAHS-SP 141
           |||||||||||||||:||:||:|:|||||||:|:|||||:||||||||||||||||| ||
Db   61 KCFCCGLMLDNWKQGDSPVEKHRQFYPSCSFVQTLLSASLQSPSKNMSPVKSRFAHS-SP 119

Qy   142 TLEHSSLPSGGSYSSLSPNPLNSRAVEDISSSRTNPSYSYAMSTEEARFLTYHMWPLTFLSP 201
           |:|||||:| ||||||||:||||||||||:|||||||||||||||||||||||:|||||||
Db   120 -LER-----GGIHSNLCSSPLNSRAVEDF-SSRMDPCSYAMSTEEARFLTYSMWPLSFLSP 173

Qy   202 SELARAGFYYIGPGDRVACFACGGGKLSNWEPKDDAMSEHRRHFPNCPFLENSLETLRFSI 261
           :|||||||||||||||||| |||||||||||||||||||||||||:||||||||:||||||
Db   174 AELARAGFYYIGPGDRVAC:FACGGGKLSNWEPKDDAMSEHRRHFPHCPFLENTSETQRFSI 233

Qy   262 SNLSMQTHAARMRTFMYWPPSVPVQPEQLASAGFYYVQPEQLASAGFYYVGRNDDVKCFPCCDGGLRCWESGDD 321
           |||||||||||||||||: :|||||||||||||||||||||||||:|||||||||||||||
Db   234 SNLSMQTHSARLRTFLYWPPSVPVQPEQLASAGFYYVDRNDDVKCLCCDGGLRCWEFGDD 293

Qy   322 PWIEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLLEQLLSTSDTPGEENADP--PIIHFG 379
           |||||||||||||||||||||||||:|||:|:||||||||||||||:||||||||| :|||||
Db   294 PWIEHAKWFPRCEFLIRMKGQEFVDEIQARYPHLLEQLLEQLLSTSDTTGEENADPTETVVHFG 353

Qy   380 PGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTTGENYKTVANDIVSALLNAED 439
           || ::||:|||||||||||:|||||:||||||||||:|||||||||||||||||||||||:
Db   354 PGE--SSKDVVMMSTPVVKALEMGFSRSLVRQTVRQTVQRQILATGENYRTVANDIVSVLLNAED 412

Qy   440 EKREEEKEKQAAEMASDLLSLIRKNRMALFQQLTCVLPILDNLLKANVINKQEHDIIKQK 499
           |:|||||||| |||||||||||||||||||||||:|||||||||:|||||||||||:||:
Db   413 ERREEEKROTEEMASGDLLSLIRKNRMALFQQLTHVLPILDNLLEASVITKQEHDIIRQK 472

Qy   500 TOIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLFVDKNMKYIPTEDVSGLSLE 559
           ||||||||||||:||||||:||||||||:||:|||||:|||:|||||||||||||||||
Db   473 TOIPLQARELIDTVLVKGNANAANIFKNSLKGIDSTLYENLFVEKNMKYIPTEDVSGLSLE 532

Qy   560 EQLRRLQEBRTCKVCMDKEVSVVFIPCGHLVVCQEBCAPSLRKCPICRGIIKGTVRTFLS 618
           |||||||||||||||||:|||:|||||||||||||||||||||||||||||||||||||||
Db   533 EQLRRLQEBRTCKVCMDRVSIVFIPCGHLVVCQEBCAPSLRKCPICRGIIKGTVRTFLS 591

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;   FILING DATE:
;   CLASSIFICATION:          514
;   ATTORNEY/AGENT INFORMATION:
;   NAME:                    Brezner, David J.
;   REGISTRATION NUMBER:     24,774
;   REFERENCE/DOCKET NUMBER: A-62464/DJB
;   TELECOMMUNICATION INFORMATION:
;   TELEPHONE:               (415)781-1989
;   TELEFAX:                 (415)398-3249
; INFORMATION FOR SEQ ID NO:  4:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH:                604 amino acids
;     TYPE:                  amino acid
;     STRANDEDNESS:          single
;     TOPOLOGY:              linear
;     MOLECULE TYPE:         protein
US-08-569-749-4

Query Match             71.8%;  Score 2353;  DB 3;   Length 604;
Best Local Similarity   72.8%;  Pred. No. 1.2e-218;
Matches   440;  Conservative  71;  Mismatches  87;  Indels   6;  Gaps   5;

Qy     20  SIMEDSTILSDWTNS-NKQKMKYDFSCELYRMSTYSTFFAGVPVSERSLARAGFYYTGVN   78
           :|:|:|            ||||  |||||  |||:|||||| ||||||||||:|||||||
Db      2  NIVENSIFLSNLMKSANTFELKYDLSCELYRMSTYSTFFAGVPVSERSLARAGFYYTGVN   61

Qy     79  DKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSA-SLGSTSKNTSP--MRNSF  135
           |||||||||||||||| | ||  || ||||| : |  :||| : |: ||:||    :||
Db     62  DKVKCFCCGLMLDNWKRGDSPTEKHKKLLYPSCRFVQSLNSVNNLEATSQPTFPSSVTNS- 120

Qy    136  AHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYAMSTEEARFLTYHMWP  195
           :||||:   ||  | ||||: | |:||:      :|| |   || || : || |||
Db    121  THSLLPGTENSGYFRGSYSNSPSNPVNSRANQDFSALMRSSYHCAMNENARLLTFQTWP  180

Qy    196  LTFLSPSELARAGFYYIGPGDRVACFACGGGKLSNWEPKDDAMSEHRRHFPNCPFLENSL-  254
           ||||||  |||||||||||||||  ||||||||||| |||| :||:||||||||||:|||
Db    181  LTFLSPTDLAKAGFYYIGPGDRVACFACGGGKLSNWEPKDNAMSEHLRHFPKCPFIENQLQ  240

Qy    255  ETLRFSISNLSMQTHAAARMRTFMYWPSSVPVQPEQLASAGFYYVGRNDDVKCFCCDGGLR  314
           :|  | ||||||||||  |:| | ||  ||| |||||||||||| | ||:|||||||||
Db    241  DTSRYTVSNLSMQTHAARFKTFFNWPSSVLVNPEQLASAGFYYVGNSDDVKCFCCDGGLR  300

Qy    315  CWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLLSTSDITTGEENADPP  374
           |||||||||| |||||||||| |:| ||| : :: ||||||||||:|:|||:
Db    301  CWESGDDPWVQHAKWFPRCEYLIRIKGQEFIRQVQASYPHLLEQLLSTSDSPGDENAESS  360

Qy    375  IIHFGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVCSKILTTGENYKTVNDIVSAL  434
           |||| |||||||  :|||||| :| | |:|||:||||| :|||||||||| |||||:||
Db    361  IIHFBPGEDHSEDAIMMNTPVINAAVEMGFSRSLVKQTVQRKILATGENYRLVNDLVLDL  420

Qy    435  LNAEDEKREBEKEKQAEEMASDDLSLIRKNRMALFQQLICVLPILDNLLKANVINKQEHD  494
           |||||:|:| :|: |: | | | |: ||||:|||| |: ||:: |||||| ||::|||||
Db    421  LNAEDEIREBERERATEBEKESNDLLIRKNRMALFQHITCVIPILDSLLTAGIINEQEHD  480

Qy    495  IIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLFVDKNMKYIPTEDVS  554
           :|||||:|| || |||:|:|| ||:||  :: |: | | |||:|||| |:|||:|||||
Db    481  VIKQKTQTSLQARELIDTILVKGNIAATVFRNSLQEAERAVLYEHLFVQQDIKYIPTEDVS  540

Qy    555  GLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLRKCPICRGIIKGTVR  614
           | ||||||||||||||||||||||:|:|||||||||||| ||||||||||| |||||||
Db    541  DLPVEEQLRRLQEERTCKVCMDKEVSIVFIPCGHLVVCKDCAPSLRKCPICRSTIKGTVR  600

Qy    615  TFLS  618
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Db    601  TFLS  604
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OM protein - protein search, using sw model
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GenCore version 5.1.6

Run on: December 4, 2003, 16:55:06 ; Search time 90.9369 Seconds
(without alignments)
1263.930 Million cell updates/sec

Title:
Perfect score: 3277
Sequence: 1 MHKTASQRLFPGPSYQNIKS.........LRKCPICRGIIKGTVRTPLS 618

Run on: US-08-569-749-2

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 684280 seqs, 185983659 residues

Total number of hits satisfying chosen parameters: 684280

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :
Published Applications AA:*
1: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
2: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
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18: /cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB ID	ID	Description
1	3277	100.0	618	12	US-10-232-286-2	Sequence 2, Appli
2	3277	100.0	618	15	US-10-153-668-338	Sequence 338, App
3	3277	100.0	618	15	US-10-207-655-200	Sequence 200, App
4	3247	99.1	618	10	US-09-974-592-8	Sequence 8, Appli
5	3247	99.1	618	12	US-10-232-286-14	Sequence 14, Appl
6	2728	83.2	612	12	US-10-232-286-14	Sequence 14, Appl
7	2724	83.1	612	10	US-09-974-592-14	Sequence 14, Appl
8	2654	81.0	591	10	US-09-201-936-42	Sequence 42, Appl
9	2353	71.8	604	12	US-10-232-286-4	Sequence 4, Appli
10	2353	71.8	604	10	US-09-201-936-13	Sequence 13, Appl
11	2332	71.2	604	12	US-10-141-618-6	Sequence 6, Appli
12	2332	71.2	604	10	US-09-974-592-6	Sequence 6, Appli
13	2326	71.0	438	8	US-10-323-643-2	Sequence 2, Appli
14	2326	71.0	438	15	US-08-464-588-2	Sequence 2, Appli
15	2172	66.3	600	10	US-09-974-592-12	Sequence 12, Appl

ALIGNMENTS

No.	Score	%	Length	DB	ID	Description
16	2152	65.7	602	10	US-09-201-936-40	Sequence 40, Appl
17	1678	51.2	1140	12	US-10-353-461-8	Sequence 8, Appl
18	1593	48.6	306	9	US-09-778-927A-62	Sequence 62, Appli
19	909	27.7	497	10	US-09-974-592-4	Sequence 4, Appli
20	909	27.7	497	10	US-09-201-936-4	Sequence 4, Appli
21	874	26.7	496	10	US-09-974-592-10	Sequence 10, Appl
22	874	26.7	496	10	US-09-201-936-10	Sequence 10, Appl
23	736.5	22.5	498	10	US-09-201-936-13	Sequence 13, Appl
24	539	16.4	108	10	US-10-228-897-7	Sequence 7, Appli
25	533.5	16.3	278	10	US-09-964-899-39	Sequence 39, Appl
26	513	15.7	268	10	US-10-041-859-2	Sequence 2, Appli
27	500.5	15.3	346	10	US-10-323-643-10	Sequence 10, Appl
28	462.5	14.1	280	15	US-08-913-322-24	Sequence 24, Appl
29	462	14.1	1403	8	US-08-913-322-22	Sequence 22, Appl
30	462	14.1	1403	15	US-10-285-408-1	Sequence 1, Appli
31	462	14.1	1403	15	US-09-201-936-12	Sequence 12, Appl
32	461	14.1	275	10	US-10-323-643-9	Sequence 9, Appli
33	461	14.1	298	15	US-10-235-026-2	Sequence 2, Appli
34	460.5	14.1	298	12	US-10-203-708-44	Sequence 44, Appli
35	457	13.9	355	12	US-10-203-708-44	Sequence 44, Appli
36	453	13.8	92	12	US-10-314-506-29	Sequence 29, Appl
37	453	13.8	92	14	US-10-014-569-29	Sequence 29, Appl
38	453	13.8	92	14	US-10-002-974-29	Sequence 29, Appl
39	453	13.8	89	9	US-09-728-721-32	Sequence 32, Appl
40	431	13.2	89	14	US-10-105-931-32	Sequence 32, Appl
41	431	13.2	89	14	US-10-118-984-32	Sequence 32, Appl
42	431	13.2	89	15	US-10-295-681-32	Sequence 32, Appl
43	419.5	12.8	172	15	US-10-041-859-8	Sequence 8, Appli
44	418.5	12.8	172	15	US-10-041-859-12	Sequence 12, Appl
45	395.5	12.1	172	15	US-10-041-859-10	Sequence 10, Appl

RESULT 1
US-10-232-286-2
Sequence 2, Application US/10232286
Publication No. US20030143579A1
GENERAL INFORMATION:
APPLICANT: Goeddel, David V
APPLICANT: Rothe, Mike
TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/232,286
FILING DATE: 30-Aug-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/569,749
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Brezner, David J.
REGISTRATION NUMBER: 24,774
REFERENCE/DOCKET NUMBER: A-62464/DJB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415)781-1989
TELEFAX: (415)398-3249
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 618 amino acids

; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-10-232-286-2

Qy   Query Match   100.0%; Score 3277; DB 12; Length 618;
            Best Local Similarity 100.0%; Pred. No. 6e-281;
Db   Matches 618; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1   MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1   MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60

Qy   61   PVSERSLARAGFYYTGVNDKVKCFCCGGLMLDNWKLGDSEIQKHKQLYPSCSFIQNLVSAS 120
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61   PVSERSLARAGFYYTGVNDKVKCFCCGGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy   121   LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121   LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180

Qy   181   MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181   MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy   241   RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241   RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy   301   NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
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Db   301   NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy   361   STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
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Db   361   STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420

Qy   421   GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD 480
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Db   421   GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD 480

Qy   481   NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 540
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   481   NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 540

Qy   541   VDKQNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
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Db   541   VDKQNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy   601   KCPICRGIIKGTVRTFLS 618
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Db   601   KCPICRGIIKGTVRTFLS 618

RESULT 2
US-10-153-668-338
; Sequence 338, Application US/10153668
; Publication No. US20030092616A1
; CURRENT FILING DATE: 2002-05-24
; GENERAL INFORMATION:
;   APPLICANT: HONDA, Goichi
;   APPLICANT: MATSUDA, Akio
;   APPLICANT: MIRAMATSU, Shuji
;   APPLICANT: ISHIZAWA, Kenya
;   TITLE OF INVENTION: STAT6 Activating Gene
;   FILE REFERENCE: 1254-0207P
;   CURRENT APPLICATION NUMBER: US/10/153,668
;   CURRENT FILING DATE: 2002-05-24
;   PRIOR APPLICATION NUMBER: US 60/293,172
;   PRIOR FILING DATE: 2001-05-25
;   PRIOR APPLICATION NUMBER: US 60/316,031
;   PRIOR FILING DATE: 2001-08-31

; PRIOR APPLICATION NUMBER: US 60/328,403
; PRIOR FILING DATE: 2001-10-12
; PRIOR APPLICATION NUMBER: JP 2001-157043
; PRIOR FILING DATE: 2001-05-25
; PRIOR APPLICATION NUMBER: JP 2001-260681
; PRIOR FILING DATE: 2001-08-30
; PRIOR APPLICATION NUMBER: JP 2001-313175
; PRIOR FILING DATE: 2001-10-10
; NUMBER OF SEQ ID NOS: 488
; SOFTWARE: Patentin Ver. 2.0
;
; SEQ ID NO 338
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-153-668-338

Qy   Query Match   100.0%; Score 3277; DB 15; Length 618;
            Best Local Similarity 100.0%; Pred. No. 6e-281;
Db   Matches 618; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1   MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1   MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60

Qy   61   PVSERSLARAGFYYTGVNDKVKCFCCGGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61   PVSERSLARAGFYYTGVNDKVKCFCCGGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy   121   LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121   LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180

Qy   181   MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181   MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy   241   RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241   RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy   301   NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   301   NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy   361   STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   361   STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420

Qy   421   GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD 480
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   421   GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD 480

Qy   481   NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 540
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   481   NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 540

Qy   541   VDKQNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
            ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   541   VDKQNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy   601   KCPICRGIIKGTVRTFLS 618
            ||||||||||||||||||
Db   601   KCPICRGIIKGTVRTFLS 618

RESULT 3
US-10-207-655-200
; Sequence 200, Application US/10207655
; Publication No. US20030118592A1
; GENERAL INFORMATION:
;   APPLICANT: Ledbetter, Jeffrey A.
;   APPLICANT: Hayden-Ledbetter, Martha S.
;   TITLE OF INVENTION: BINDING DOMAIN-IMMUNOGLOBULIN FUSION PROTEINS

; FILE REFERENCE: 390069.401C1
; CURRENT APPLICATION NUMBER: US/10/207,655
; CURRENT FILING DATE: 2002-07-25
; NUMBER OF SEQ ID NOS: 426
; SOFTWARE: Patentin version 3.0
;
; SEQ ID NO 200
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-207-655-200

Query Match   100.0%; Score 3277; DB 15; Length 618;
Best Local Similarity   100.0%; Pred. No. 6e-281;
  Matches 618; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMYDFSCELYRMSTYSTFPAGV 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMYDFSCELYRMSTYSTFPAGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy   121 LGSTSKNTSPMRNSPAHSLSPTLEHSSLFSGSYSSLSPNELNSRAVEDISSSRTNPYSYA 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121 LGSTSKNTSPMRNSPAHSLSPTLEHSSLFSGSYSSLSPNELNSRAVEDISSSRTNPYSYA 180

Qy   181 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy   301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy   361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420

Qy   421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIXRKNRMALFQQLTCVLPILD 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIXRKNRMALFQQLTCVLPILD 480

Qy   481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIIVKGNAAANIFKNCLKEIDSTLYXNLF 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIIVKGNAAANIFKNCLKEIDSTLYXNLF 540

Qy   541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy   601 KCPICRGIIKGTVRTFLS 618
              ||||||||||||||||||
Db   601 KCPICRGIIKGTVRTFLS 618

RESULT 4
US-09-974-592-8
Sequence 8, Application US/09974592
Patent No. US20020120121A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE

; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009004
; CURRENT APPLICATION NUMBER: US/09/974,592
; CURRENT FILING DATE: 2001-10-09
; PRIOR APPLICATION NUMBER: US 09/617,053
; PRIOR FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 08/800,929
; PRIOR FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSEQ for Windows Version 4.0
;
; SEQ ID NO 8
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-974-592-8

Query Match   99.1%; Score 3247; DB 10; Length 618;
Best Local Similarity   99.4%; Pred. No. 2.7e-278;
  Matches 614; Conservative   0; Mismatches   4; Indels   0; Gaps   0;

Qy   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMYDFSCELYRMSTYSTFPAGV 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1 MHKTASQRLFPGPSYQNIKSIMEDSTILSDWTNSNKQKMYDFSCELYRMSTYSTFPAGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy   121 LGSTSKNTSPMRNSPAHSLSPTLEHSSLFSGSYSSLSPNELNSRAVEDISSSRTNPYSYA 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121 LGSTSKNTSPMRNSPAHSLSPTLEHSSLFSGSYSSLSPNELNSRAVEDISSSRTNPYSYA 180

Qy   181 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy   301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy   361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVLSKILTT 420

Qy   421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIXRKNRMALFQQLTCVLPILD 480
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIXRKNRMALFQQLTCVLPILD 480

Qy   481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIIVKGNAAANIFKNCLKEIDSTLYXNLF 540
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIIVKGNAAANIFKNCLKEIDSTLYXNLF 540

Qy   541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
              |||||||||||||||||||||||| |||||||||||||||||||||||||||||||||||
Db   541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy   601 KCPICRGIIKGTVRTFLS 618
              ||||||||||||||||||
Db   601 KCPICRGIIKGTVRTFLS 618

RESULT 5
US-09-201-936-8
Sequence 8, Application US/09201936
Publication No. US20020187946A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: MacKenzie, Alexander B.
; APPLICANT: Baird, Stephen

RESULT 6
US-10-232-286-14

; APPLICANT: Liston, Peter
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; FILE REFERENCE: 07891/003003
; CURRENT APPLICATION NUMBER: US/09/201,936
; CURRENT FILING DATE: 1998-12-01
; EARLIER APPLICATION NUMBER: 09/011,356
; EARLIER FILING DATE: 1998-02-04
; EARLIER APPLICATION NUMBER: PCT/IB96/01022
; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 08/576,956
; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 1995-12-22
; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSEQ for Windows Version 3.0
US-09-201-936-8
; SEQ ID NO 8
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens

Query Match   99.1%;
Best Local Similarity   99.4%; Score 3247; DB 10; Length 618;
Matches 614; Conservative 0; Mismatches 4; Indels 0; Gaps 0; Pred. No. 2.7e-278;

Qy   1 MHKTASQRLPFGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1 MHKTASQRLPFGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSAS 120

Qy 121 LGSTSKNTSPMRNSPAHSLSPTLEHSSLFSGGYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLLPPNPLNSRAVEDISSSRTNPYSYA 180

Qy 181 MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVACFACCGGKLSNWEPKDDAMSEH 240
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 181 MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240

Qy 241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300

Qy 301 NDDVCKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 301 NDDVCKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy 361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVLSKILTT 420

Qy 421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQOLTCVLPILD 480
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQOLTCVLPILD 480

Qy 481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAA.ANIFKNCLKEIDSTLYKNLF 540
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTIWVKGNAA.ANIFKNCLKBIDSTLYKNLF 540

Qy 541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600

Qy 601 KCPICRGIIKGTVRTFLS 618
        ||||||||||||||||||
Db 601 KCPICRGIIKGTVRTFLS 618

; Sequence 14, Application US/10232286
; Publication No. US20030143579A1
; GENERAL INFORMATION:
; APPLICANT: Roethe, Mike
; APPLICANT: Goeddel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/232,286
; FILING DATE: 30-Aug-2002
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749
; FILING DATE: <Unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: Brezner, David J.
; REGISTRATION NUMBER: 24,774
; REFERENCE/DOCKET NUMBER: A-62464/DJB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415)781-1989
; TELEFAX: (415)398-3249
; INFORMATION FOR SEQ ID NO: 14:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 612 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-10-232-286-14

Query Match   83.2%;
Best Local Similarity   83.4%; Score 2728; DB 12; Length 612;
Matches 517; Conservative 48; Mismatches 45; Indels 10; Gaps 6; Pred. No. 2.3e-232;

Qy   1 MHKTASQRLPFGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60
        |||| :||||| :|| : |||||| |||| || :| |||||||||||||||| : | |
Db   1 MDKTVSQRLQGQTLHQKLKRIMEKSTILSNWTKRSBEKMKPDFSCELYRMSTYSAPPRGV 60

Qy   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKQGDSPVEKHRQFYPSCSFVQTLLSAS 120
        |||||||||||||||||||||||||||||||||| |||| : |:: |||||| : | |
Db   61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKQGDSPVEKHRQFYPSCSFVQTLLSAS 120

Qy 121 LGSTSKNTSPMRNSFAHSLSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
        ||||||||| ||:||:||:| :|| | |||||:: |:: |:| | || :
Db 121 LQSPSKNMSPVKSRFAHS-SP-LER----GGIHSNLCSSPLNSRAVEDF-SSRMDPCSYA 173

Qy 181 MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVACFACCGGKLSNWEPKDDAMSEH 240
        |||||||| :|::|| :||:|:|||||||||||||||||||| ||||||||||||||||||
Db 174 MSTEEARFLTYSMWPLSPFLSPABLARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 233

Qy 241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
        |||||| |||| :| |||||||||:||| || : | : ||||||||||||||||||||: |
Db 234 RRHFPHCPFLENTSETQRFSISNLSMQTHSARLRTFLYWPPSVPVQPEQLASAGFYYVDR 293

Qy 301 NDDVCKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
        |||||||||||||||||||:|||||||||||||||||||||||||||||||||||||||||
Db 294 NDDVCKCFCCDGGLRCWEPGDDPWIEHAKWFPRCEFLIRMKGQEFVDEIQARYPHLLEQLL 353

Qy 361 STSDTTGEENADPPIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILT 418
        |||||||||||||--||||||||||||||||||||||||||||||||||||||| :| |
Db 361 STSDTTGEENADP--PIIHFGPGESSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKIL 418

RESULT 7
US-09-974-592-14

; Sequence 14, Application US/09974592
; Patent No. US20020121A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009004
; CURRENT APPLICATION NUMBER: US/09/974,592
; CURRENT FILING DATE: 2001-10-09
; PRIOR APPLICATION NUMBER: US 09/617,053
; PRIOR FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 08/800,929
; PRIOR FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SEQ ID NO 14
; LENGTH: 612
; TYPE: PRT
; ORGANISM: Mus musculus
US-09-974-592-14

Query Match   83.1%;   Score 2724;   DB 10;   Length 612;
Best Local Similarity   83.2%;
Matches 516;   Conservative 46;   Mismatches 48;   Indels 10;   Gaps 6;

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Qy      1 MHKTASQRLFPPGPSYQNIKSIMEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGV 60
            |||||||||||||||||||:||||||||||||||||||||| ::||||||||||||||||
Db      1 MDKTVSQRLGQGTLHQKLRIMEKSTILSNWTKESEKMKFDFSCELYRMSTYSAFPRGV 60

Qy     61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKQGDSPVEKHRQFYPSCSFVQTLLSAS 120
            ||||||||||||||||||||||||||||||||||:|||||||||||||||||||:||||
Db     61 PVSERSLARAGFYYTGVNDKVKCFCCGLMLDNWKQGDSPIQHKQLYPSCSFIQNLVSAS 120

Qy    121 LGSTSKNTSPMRNSPAHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYA 180
            |||||||||||||||||::||||||| |||| |:|||||||||||||||:|| ||||||
Db    121 LQSPSKNMSPVKSRPAHS-SP-LER----GGIHSNLCSSPLNSRAVEDF-SSRNDPCSYA 173

Qy    181 MSTEEARFLITYHNWPLTFLSPSELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 240
            |||||||||||||||||||||:|||:|||||||||||||||||||||||||||||||||
Db    174 MSTEEARFLITYSNWPLSFLSPABLARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEH 233

Qy    241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
            ||| ||||||||||||||| :|:|||:||:|||:|||||||: :||:|||||||||||
Db    234 RRHFPHCPFLENTSETQRFSISNLSMQTHSARLRTFLYWPPSSVPVQPEQLASAGFYYDR 293

Qy    301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
            |||||||||||||||||||||||||||||||||||||||||||:||||||||||||||
Db    301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360

Qy    361 STSDTPGEENADPTETVVHFGPGE-SSEDVVMMSTPVVKAALEMGFSRSLVRQTVQRQIL 412
            :||:||||||:|-PIIHFGPGESSSEDAVMMNIPVVKSALEMGFNRDLVKQTVQSKIL
Db    354 STSDTPGEENAD---PIIHFGPGESSSEDAVMMNIPVVKSALEMGFNRDLVKQTVQSKIL 412

Qy    413 ATGENYRITVNDIVSVLLNAEDERREEEKERQTEEMASGDJSLIRKNRMALFQQLTHVLPI 472
            ||:|||||||||||||||||||:|:| |||:||||||:||::|||||||||||| |||
Db    413 ATGENYRITVNDIVSVLLNAEDERREEEKERQTEEMASGDLSLIRKNRMALFQQLTHVLPI 472

Qy    473 LDNLLEASVTTKQEHDIIRQKTQIPLQARELIDTVLVKGNAAANIFKNLKEIDSTLYKN 538
            |||||||||||:|||||:||||||||||||||||:||||||||||||:||||||||||
Db    473 LDNLLEASVTTKQEHDIIRQKTQIPLQARELIDTVLVKGNAAANIFKNLKEIDSTLYKN 538

Qy    539 LFVDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQBCAPS 598
            |||||||||||||||||||||||||||:||| ||:|||||:|||||||||||||||||
Db    539 LFVEKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDREVSIVFIPCGHLVVCQBCAPS 592

Qy    599 LRKCPICRGIIKGTVRTFLS 618
            |||||||||||||||||||
Db    593 LRKCPICRGTIKGTVRTFLS 612
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RESULT 8
US-09-201-936-42

; Sequence 42, Application US/09201936
; Publication No. US20020187946A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: MacKenzie, Alexander E.
; APPLICANT: Baird, Stephen
; APPLICANT: Liston, Peter
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; FILE REFERENCE: 07891/003003
; CURRENT APPLICATION NUMBER: US/09/201,936
; CURRENT FILING DATE: 1998-12-01
; EARLIER APPLICATION NUMBER: 09/011,356
; EARLIER FILING DATE: 1998-02-04
; EARLIER APPLICATION NUMBER: PCT/IB96/01022
; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 08/576,956
; EARLIER FILING DATE: 1995-12-22
; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 42
; LENGTH: 591
; TYPE: PRT
; ORGANISM: Mus musculus
US-09-201-936-42

Query Match   81.0%;   Score 2654;   DB 10;   Length 591;
Best Local Similarity   83.8%;
Matches 502;   Conservative 44;   Mismatches 43;   Indels 10;   Gaps 6;

```
Qy     22 MEDSTILSDWTNSNKQKMKYDFSCELYRMSTYSTFPAGVPVSERSLARAGFYYTGVNDKV 81
            |||||||||||||||||||| ::||||||||||||||||||||||||||||||||||||
Db      1 MEKSTILSNWTKESEKMKFDFSCELYRMSTYSAFPRGVPVSERSLARAGFYYTGVNDKV 60

Qy     82 KCFCCGLMLDNWKQGDSPIQHKQLYPSCSFIQNLVSASLGSTSKNTSPMRNSPAHSLSP 141
            ||||||||||||||||||:|||||||||||:||||:||| ||||||||||||||||||
Db     61 KCFCCGLMLDNWKQGDSPVEKHRQFYPSCSFVQTLLSAS-LQSPSKNMSPVKSRPAHS-SP 119

Qy    142 TLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYAMSTEEARFLITYHNWPLTFLSP 201
            ||||| |||| |:|||||||||||||:|| |||||||||||||||||||||||:||||
Db    120 -LER----GGIHSNLCSSPLNSRAVEDF-SSRMDPCSYAMSTEEARFLITYSNWPLSFLSP 173

Qy    202 SELARAGFYYIGPGDRVACFACGGKLSNWEPKDDAMSEHRRHFPNCPFLENSLETLRFSI 261
            ::|||||||||||||||||||||||||||||||||||||:||| ||||||||||||||
Db    174 AELARAGFYYTGPGDRVACFACGGKLSNWEPKDVAMSEHRRHFPHCPFLENTSETQRFSI 233
```

RESULT 9
US-10-232-286-4
Sequence 4, Application US/10232286
Publication No. US20030143579A1
GENERAL INFORMATION:

APPLICANT: Roche, Mike
        Goeddel, David V

TITLE OF INVENTION: INHIBITORS OF APOPTOSIS

NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/232,286
FILING DATE: 30-Aug-2002
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/569,749
FILING DATE: <Unknown>

ATTORNEY/AGENT INFORMATION:
NAME: Brezner, David J.
REGISTRATION NUMBER: 24,774
REFERENCE/DOCKET NUMBER: A-62464/DJB

TELECOMMUNICATION INFORMATION:
TELEPHONE: (415)781-1989
TELEFAX: (415)398-3249

INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 604 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 4:

US-10-232-286-4

Query Match   71.8%; Score 2353; DB 12; Length 604;
Best Local Similarity   72.8%; Pred. No. 3.5e-199;

RESULT 10
US-10-141-618-6
Sequence 6, Application US/10141618
Publication No. US20030165887A1
GENERAL INFORMATION:

APPLICANT: Reed, John C.

TITLE OF INVENTION: Methods For Determining the Prognosis
    For Cancer Patients Using Tucan

FILE REFERENCE: P-LJ 5254

CURRENT APPLICATION NUMBER: US/10/141,618
CURRENT FILING DATE: 2002-05-07

PRIOR APPLICATION NUMBER: US 60/356,934
PRIOR FILING DATE: 2002-02-12

PRIOR APPLICATION NUMBER: US 09/388,221
PRIOR FILING DATE: 1999-09-01

NUMBER OF SEQ ID NOS: 15

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 6
LENGTH: 604
TYPE: PRT
ORGANISM: Homo sapiens

US-10-141-618-6

Query Match   71.8%; Score 2353; DB 12; Length 604;

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; SEQ ID NO 6
;    LENGTH: 604
; SOFTWARE: FastSEQ for Windows Version 4.0
;    NUMBER OF SEQ ID NOS: 17
;    PRIOR FILING DATE: 1997-02-13
;    PRIOR APPLICATION NUMBER: US 08/800,929
;    PRIOR FILING DATE: 2000-07-14
;    PRIOR APPLICATION NUMBER: US 09/617,053
;    CURRENT FILING DATE: 2001-10-09
;    CURRENT APPLICATION NUMBER: US/09/974,592
;    FILE REFERENCE: 07891/009004
; TITLE OF INVENTION: DISEASE
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
;    APPLICANT: Pratt, Christine
;    APPLICANT: Tsang, Benjamin K
;    APPLICANT: Baird, Stephen
;    APPLICANT: Liston, Peter
;    APPLICANT: MacKenzie, Alexander E
;    APPLICANT: Korneluk, Robert G
; GENERAL INFORMATION:
; Patent No. US20020120121A1
; Sequence No. 6, Application US/09974592
US-09-974-592-6

RESULT 11

Qy    20 SIMEDSTILSDWTNS-NKQKMKYDFSCELYRMSTYSTPPNGVPVPVSBRSLARAGFYYTGVN  78
         :|:|:| || ||  :::|||.||||||||||||||||||||||||||||||||||||||||
Db     2 NIVENSIFLSNLMKSANTFELKYDLSCELYRMSTYSTPPAGVPVPVSERSLARAGFYYTGVN  61

Qy    79 DKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSA--SLGSTSKNTSP--MRNSF 135
         ||||||||||:|||||||||:||||||||||||||||||||  :||||||||||  ||||
Db    62 DKVKCFCCGLMLDNWKRGDSPTEKHKKLYPSCRFVQSLNSVNNLEATSQPTFPSSVTNS- 120

Qy   136 AHSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYAMSTEEARFLTYHMWP 195
         || |||||:|| ::||  | : |||| ::||||||| ||:|:|| ||::||||| |||||
Db   121 THSLLPGTENSGYFRGSYSNSPSNPVNSRANQDFSALMRSSYHCAMNNENARLLTFQTWP 180

Qy   196 LTFLSPSELARAGFYYIGPGDRVACFACGGGKLSNWEPKDDAMSEHRRHFPNCPFLENSL- 254
         :||:|| :|||||||:|||||||||||||||:|||||:|:::||||||||| ||||||
Db   181 LTFLSPTDLAKAGFYYIGPGDRVACFACGGGKLSNWEPKDNAMSEHLRHFPKCPFIENQLQ 240

Qy   255 ETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPBQLASAGFYYVGRNDDVKCFCCDGGLR 314
         : |||||:||||||||||||||:|||| ||||||||| | |:||||||||||||||||
Db   241 DTSRYTVSNLSMQTHAARFKIFPFNWPSSVLVNPEQLASAGFYYVGNSDDVKCFCCDGGLR 300

Qy   315 CWESGDDPWVEHAKWFPRCEFLIRMKQOEFVDEIQGRYPHLLEQLLSTSDTTGBENADPP 374
         |||||||||||:||||||||||||||||::|:|:|||||||:|||||:|:|| :|:||
Db   301 CWESGDDPWVQHAKWFPRCEYLIRIKGQEFIRQVQASYPHLLEQLLSTSDSPGDENAESS 360

Qy   375 IIHFGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTTGENYKTVNDIVSAL 434
         ||||||:|||||||:||||||||||||||||||||:|||||:|||||||:||||||||:
Db   361 IIHFPEGEDHSEDAIMMNTPVINAAVEMGFSRSLVKQTVQRKILATGENYRLVNDLVLDL 420

Qy   435 LNAEDEKREEEKEKQOAEEMASDDLSLIRKNRMALFQOLTCVLPILIDNLLKANVINKQEHD 494
         |||||| |||||||||||||||||||||||||||||||||||||||:||||||||||||||
Db   421 LNAEDEIREEERRATEEKESNDLLLIRKNRMALFQHLTCVIPILDSLLTAGIINEQEHD 480

Qy   495 IIKQKTQIPLQARELIDTILVKQNAAANIFKNCLKEIDSTLYKNLFVDKNMKYIPTEDVS 554
         :||||||||||||||||||:|||||||||||||||||||||:|:|||||||||||||:
Db   481 VIKQKTQTSLQARELIDTILVKGNIAATVFRNSLQEAEAVLYEHLFVQQDIKYIPTEDVS 540

Qy   555 GLSLEBEQLRRLQEBRTCKVCMDKEVSVVFIPCGHLVVCQBCAPSLRKCPICRGIIKGTVR 614
         |||:||||||||||||||||||||||||||||||||:||:||||||||||||||:||||
Db   541 DLPVEBQLRRLQEBRTCKVCMDKEVSIVFIPCGHLVVCKDCAPSLRKCPICRSTIKGTVR 600

Qy   615 TFLS 618
         ||||
Db   601 TFLS 604
```

```
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-974-592-6

Best Local Similarity  72.8%;    Pred. No. 3.5e-199;    Length  604;
Matches 440; Conservative  71;    Mismatches  87;    Indels   6;    Gaps   5;

RESULT 12

Qy    20 SIMEDSTILSDWTNS-NKQKMKYDFSCELYRMSTYSTPPNGVPVPVSBRSLARAGFYYTGVN  78
         :|:|:| || ||  :::|||.||||||||||||||||||||||||||||||||||||||||
Db     2 NIVENSIFLSNLMKSANTFELKYDLSCELYRMSTYSTPPAGVPVPVSERSLARAGFYYTGVN  61

Qy    79 DKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSA--SLGSTSKNTSPMRNSF 136
         ||||||||||:|||||||||:||||||||||||||||||||  :||||||||||||||
Db     2 DKVKCFCCGLMLDNWKRGDSPTEKHKKLYPSCRFVQSLNSVNNLEATSQPTFPSSVTHST 121

Qy   137 HSLSPTLEHSSLFSGGSYSSLSPNPLNSRAVEDISSSRTNPYSYAMSTEEARFLTYHMWPL 196
         ||:|||||:|| ::||  | : |||| ::||||||| ||:|:|| ||::||||| |||||
Db   122 HSLLPGTENSGYFRGSYSNSPSNPVNSRANQBFSALMRSSYPCMNNENARLLTFQTWPL 181

Qy   197 TFLSPSELARAGFYYIGPGDRVACFACGGGKLSNWEPKDDAMSEHRRHFPNCPFLENSL-E 255
         |||:|| :|||||||:|||||||||||||||:|||||:|:::||||||||| ||||||
Db   182 TFLSPTDLARAGFYYIGPGDRVACFACGGGKLSNWEPKDNAMSEHLRHFPKCPFIENQLQD 241

Qy   256 TLRFSISNLSMQTHAARMRTFMYWPSSVPVQPBQLASAGFYYVGRNDDVKCFCCDGGLRC 315
         | |||||:||||||||||||||:|||| ||||||||| | |:||||||||||||||||
Db   242 TSRYTVSNLSMQTHAARKTFPFNWPSSVLVNPEQLASAGFYYVGNSDDVKCFCCDGGLRC 301

Qy   316 WESGDDPWVEHAKWFPRCEFLIRMKQOEFVDEIQGRYPHLLEQLLSTSDTTGBENADPPI 375
         |||||||||||:||||||||||||||||::|:|:|||||||:|||||:|:|| :|:||
Db   302 WESGDDPWVQHAKWFPRCEYLIRIKGQEFIRQVQASYPHLLEQLLSTSDSPGDENABSSI 361

Qy   376 IHFGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTTGENYKTVNDIVSALL 435
         |||||:|||||||:||||||||||||||||||||:|||||:|||||||:||||||||:
Db   362 IHLPEGEDHSEDAIMMNTPVINAAVEMGFSRSLVKQTVQRKILATGENYRLVNDLVLDLL 421

Qy   436 NAEDEKREEEKEKQOAEEMASDDLSLIRKNRMALFQOLTCVLPILIDNLLKANVINKQEHDI 495
         |||||| |||||||||||||||||||||||||||||||||||||||:||||||||||||||
Db   422 NAEDEIREEERRATEEKESNDLLLIRKNRMALFQHLTCVIPILDSLLTAGIINEQEHDV 481

Qy   496 IKQKTQIPLQARELIDTILVKQNAAANIFKNCLKEIDSTLYKNLFVDKNMKYIPTEDVSG 555
         :||||||||||||||||||:|||||||||||||||||||||:|:|||||||||||||:
Db   482 IKQKTQTSLQARELIDTILVKGNIAATVFRNSLQEAEAVLYEHLFVQQDIKYIPTEDVSD 541

Qy   556 LSLEBEQLRRLQEBRTCKVCMDKEVSVVFIPCGHLVVCQBCAPSLRKCPICRGIIKGTVRT 615
         |||:||||||||||||||||||||||||||||||||:||:||||||||||||||:||||
Db   542 LPVEBEQLRRLPEBRTCKVCMDKEVSIVFIPCGHLVVCKDCAPSLRKCPICRSTIKGTVRT 601

Qy   616 FLS 618
         |||
Db   602 FLS 604
```

```
; Query Match  435;    Conservative  72;    Mismatches  92;    Indels   4;    Gaps   4;
Best Local Similarity  72.1%;    Pred. No. 2.5e-197;
Matches 435;    Score 2332;    DB 10;    Length 604;

US-09-201-936-6

RESULT 12

;    EARLIER FILING DATE: 1995-12-22
;    EARLIER APPLICATION NUMBER: 08/576,956
;    EARLIER FILING DATE: 1996-08-05
;    EARLIER APPLICATION NUMBER: PCT/IB96/01022
;    EARLIER FILING DATE: 1998-02-04
;    EARLIER APPLICATION NUMBER: 09/011,356
;    EARLIER FILING DATE: 1998-12-01
;    CURRENT APPLICATION NUMBER: US/09/201,936
;    CURRENT FILING DATE: 1998-12-01
;    FILE REFERENCE: 07891/003003
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
;    APPLICANT: Liston, Peter
;    APPLICANT: Baird, Stephen
;    APPLICANT: MacKenzie, Alexander E.
;    APPLICANT: Korneluk, Robert G.
; GENERAL INFORMATION:
; Publication No. US20020187946A1
; Sequence 6, Application US/09201936
US-09-201-936-6
```

; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSEQ for Windows Version 3.0
;
; SEQ ID NO 6
; LENGTH: 604
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-201-936-6

Query Match   71.2%; Score 2332; DB 10; Length 604;
Best Local Similarity   72.1%; Pred. No 2.5e-197;
Matches 435; Conservative 72; Mismatches 92; Indels 4; Gaps 4;

```
Qy   20  SIMEDSTILLSDWTNS-NKQKMKYDFSCELYRMSTYSTFFAGVPVSERSLARAGFYYTGVN   78
         |:|:|  ||:|:  |||     ||:|||||||||||||||||||||||
Db    2  NIVENSIFLSNLMKSANTFELKYDLSCELYRMSTYSTFFAGVPVSERSLARAGFYYTGVN    61

Qy   79  DKVKCFCCGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSA-SLGSTSKNTSPMRNSFA-  136
         |||||||||||    | : |||||||:|:|   |||||||  :|:|:||||   |:
Db   62  DKVKCFCCGLMLDNWKRGDSPTEKHKKLYPSCRFVQSLNSVANNLEATSQPTFPSSVTHST  121

Qy  137  HSILSPTLEHSSLFSGSYSSLSPNPLNSRAVEDISSSRTNPYSYAMSTEEARFLTYHMWPL  196
         |||    ||:   |      :||||   ::|| ||  :|:  |:|||||    |||| |:
Db  122  HSILPGTENSGYFRGSYSNSPSNPVNSRANQEFSALMRSSYPCPMNNENARLLTFQTWPL   181

Qy  197  TFPLSPSELARAGFYYIGPGDRVACFACGGGKLSNWEPKDNAMSEHLRHPPKCPFIENQLQD  241
         |||||   |||||||||||||||||||||||||||||| : |||   ::|| |:|||  ::|
Db  182  TFLSPTDLARAGFYYIGPGDRVACFACGGGKLSNWEPKDNAMSEHLRHPPKCPFIENSL-E  255

Qy  242  TSRYTVSNLSMQTHAARRTFMYWPSSVLVNPEQLASAGFYYVGNSDDVKCFCCDGGLRC   301
         |:|: |:||||||||||   ||   |||   |:||||||||||:   ||||||||||:|||
Db  256  TLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGRNDDVKCFCCDGGLRC   315

Qy  316  WESGDDPWVEHAKWFPRCEFLIRMKGQOEFVDEIQGRYPHLLEQLLSTSDTTGBENADPPI  375
         |||||||||:|||||||||:|||:||||||:| :  :|||||||||||||:  |:|||| :|
Db  302  WESGDDPWVQHAKWFPRCEYLIRIKGQOEFIRQVQASYPHLLEQLLSTSDSPGDENABSSI  361

Qy  376  IHFGPEGSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTTGENYKTVNDIVSALL   435
         || :|| . ||||:||||||:.|.|||||.|.|||||||:|||.|||||:.||||:.| |
Db  362  IHLEPGEDHSEDAIMMNTPVINAAVEMGFSRSLVKQTVQRKILATGENYRLVNDLVLDLI   421

Qy  436  NABDEKREBEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILDNILIKANVINKQEHDI  495
         ||.|| |.||| .| |:.|||:||||||||||||||:|||:.|:||.:|:.|||:|:|||:
Db  422  NAEDEIREEEBRATEBKEBSNDLLLIRKNRMALFQHLTCVIPLLDSLLITAGIINBQEHDV  481

Qy  496  IKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYXNLFVDKNMKYIPTEDVSG   555
         |||||| |||||||||||||||.|||.:|:|.|:||.|.|||.||||.||||||||||||:
Db  482  IKQKTQTSLQARELIDTILVKGNIAATVFRNSLQEABAAVLYEHLFVQQDIKYIPTEDVSD  541

Qy  556  LSLEEQLRRLQEERTCKYCMDKEVSVVFIPCGHLVVCQECAPSLRKCPICRGIIKGTVRT  615
         | .|||||||.|||||:|||||||.:|||||||||||:.|||||||||||||:.|||||||
Db  542  LPVEEOQLRRLPEERTCKVCMDKEVSIVFIPCGHLVVCKDCAPSLRKCPICRSTIKGTVRT  601

Qy  616  FLS 618
         |||
Db  602  FLS 604
```

Query Match   71.0%; Score 2326; DB 8; Length 438;
Best Local Similarity   100.0%; Pred. No 5.1e-197;
Matches 438; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
Qy  181  MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGGKLSNWEPKDDAMSEH  240
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1  MSTEEARFLTYHMWPLTFLSPSELARAGFYYIGPGDRVACFACGGGKLSNWEPKDDAMSEH   60

Qy  241  RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR   300
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61  RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR   120

Qy  301  NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQOEFVDEIQGRYPHLLEQLL  360
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  121  NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQOEFVDEIQGRYPHLLEQLL  180

Qy  361  STSDTTGBENADPPIIHFGPGBSSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT   420
         |||||||||||||||||||||:||||||||||||||||||||||||||||||||||||||
Db  181  STSDTTGBENADPPIIHFGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT   240

Qy  421  GENYKTVNDIVSALLNABEDEKREBEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD  480
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  241  GENYKTVNDIVSALLNABEDEKREBEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD  300

Qy  481  NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYXNLF   540
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  301  NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYXNLF   360

Qy  541  VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKYCMDKEVSVVFIPCGHLVVCQECAPSLR   600
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  361  VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKYCMDKEVSVVFIPCGHLVVCQECAPSLR   420

Qy  601  KCPICRGIIKGTVRTFLS 618
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Db  421  KCPICRGIIKGTVRTFLS 438
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; TITLE OF INVENTION: Human Inhibitor of Apoptosis Gene 1
; FILE REFERENCE: PF165P1D1
; CURRENT APPLICATION NUMBER: US/10/323,643
; CURRENT FILING DATE: 2002-12-20
; PRIOR APPLICATION NUMBER: 08/464,588
; PRIOR FILING DATE: 1995-06-05
; PRIOR APPLICATION NUMBER: PCT/US95/05922
; PRIOR FILING DATE: 1995-05-11
; NUMBER OF SEQ ID NOS: 10
; SEQ ID NO 2
; LENGTH: 438
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-323-643-2
; SOFTWARE: Patentin version 3.1

Query Match      100.0%;  Score 2326;  DB 15;  Length 438;
Best Local Similarity   71.0%;  Pred. No. 5.1e-197;
Matches  438;  Mismatches   0;  Conservative   0;  Indels   0;  Gaps   0;

Qy     181 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFPACGGKLSNWEPKDDAMSEH 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1 MSTEEARFLITYHMWPLTFLSPSELARAGFYYIGPGDRVACFPACGGKLSNWEPKDDAMSEH 60

Qy     241 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 300
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      61 RRHFPNCPFLENSLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGR 120

Qy     301 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     121 NDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLL 180

Qy     361 STSDTTGEENADPPIIHFGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     181 STSDTTGEENADPPIIHFGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTT 240

Qy     421 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     241 GENYKTVNDIVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILD 300

Qy     481 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     301 NLLKANVINKQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLF 360

Qy     541 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     361 VDKNMKYIPTEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLR 420

Qy     601 KCPICRGIIKGTVRTFLS 618
           ||||||||||||||||||
Db     421 KCPICRGIIKGTVRTFLS 438
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Query Match      66.3%;  Score 2172;  DB 10;  Length 600;
Best Local Similarity   66.7%;  Pred. No. 3.6e-183;
Matches  406;  Mismatches  96;  Conservative  87;  Indels  20;  Gaps  7;

Qy      21 IMEDSTILSDWTNS-NKQKMKYDFSCELIYRMSTIYSTFPAGVPVSERSLARAGFYYTGVND 79
           :::||  | :  ||  ||: : |   |:|| ::| | | ||||:||: ||||||||||||
Db       1 MVQDSAFLAKLMKSADTFELKYDFSCELIYRLSTYSAFPRGVPVSERSLARAGFYYTGVND 60

Qy      80 KVKCFCCGGLMLDNWKLGDSPIQKHKQLYPSCSFIQNLVSA------SLGSTSKNTSP 130
           ||||||||||||||||||:|||  |  | |:|| || ::|       ::|: |: |
Db      61 KVKCFCCGGLMLDNWKGDSPMEKHRKLYPSCNFVQTLNPANSLRASPRPSLPSTAMSTMP 120

Qy     131 MRNSFAHSLSPTLEHSSLFSCGSYSSLSPNPLNSRAVEDISSSRTNPYSYAMSTEEARFLT 190
           :|  | ||||: || ||| | | || |: |  ||  |:|:  :||| |::|||||||||
Db     121 L--SFASS-----ENTGYFSGSYSSFPSDPVNFRANQCDPALSTSPYHFANNTEKARLLT 173

Qy     191 YHMWPLTFLSPSELARAGFYYIGPGDRVACFPACGGKLSNWEPKDDAMSEHRRHFPNCPFL 250
           |||||||||||||||||||||||||||||||| ||||||||:|||||||:||||||||||
Db     174 YETWPLSFLSPAKLAKAGFYYIGPGDRVACFACDGKLSNWERKDDAMSEHQRHFPSCPFL 233

Qy     251 EN-SLETLRFSISNLSMQTHAARMRTFMYWPSSVPVQPEQLASAGFYYVGRNDDVKCFCC 309
           || ||||||||||||||||||||||||||||||||||:||||||||||||:|||||||
Db     234 KDLGQSASRYTVSNLSMQTHAARIRTFSNWPSSALVHSQELASAGFYYTGHSDDVKCFCC 293

Qy     310 DGGLRCWESGDDPWVEHAKWFPRCEFLIRMKGQEFVDEIQGRYPHLLEQLLSTSDTTGEE 369
           |||||||||||||||||||||||:  ||| |::| :|| ||||:||||||||:|: | |
Db     294 DGGLRCWESGDDPWVEHAKWFPRCEYLIRIKGQEFVSQVQAGYPHLLEQLLSTSDSPBDS 353

Qy     370 NADPPIIHFGPGESSSEDAVMMNTPVVKSALEMGFNRDLVKQTVQSKILTTGENYKTVND 429
           ||:|||||||| || |||||| |||||||||:||| |:||:|| : |:||| ||||||
Db     354 NADAAIVHFGPGE-SSEDVVMMSTPVVKAALEMGFSRSLSVRQTVQRQILATGENYRTVSD 412

Qy     430 IVSALLNAEDEKREEEKEKQAEEMASDDLSLIRKNRMALFQQLTCVLPILDNLLKANVIN 489
           ||||:|| |:| |:||: :| ::|||||||||||||:| ||||  | || || |||:||
Db     413 LVIGLLDAEDEMREEQMEQAABEEESSDDLALIRKNKMVLFQHLTCVTPMLYCLLSARAIT 472

Qy     490 KQEHDIIKQKTQIPLQARELIDTILVKGNAAANIFKNCLKEIDSTLYKNLFVDKNMKYIP 549
           | | |:|| ||  | | || |:| ||  |  | | : | ||||| ||| |:|||||||
Db     473 EQECNAVRQKPH-TLQASTLIDTVLAKGNTAATSFRNSLREIDPALYRDIFVQQDIRSLD 531

Qy     550 TEDVSGLSLEEQLRRLQEERTCKVCMDKEVSVVFIPCGHLVVCQECAPSLRKCPICRGII 609
           | | ::|:||||||||||||||:|| |:|:|  ||||||||||:|||: |||||||||
Db     532 TDDIAALPMEBQLRKLQEERMCKVCMDREVSIVFIPCGHLVVCKDCAPSLRKCPICRGTI 591

Qy     610 KGTVRTFLS 618
           |||||||||
Db     592 KGTVRTFLS 600
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Search completed: December 4, 2003, 17:19:23
Job time : 92.9369 secs

OM protein - protein search, using sw model

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GenCore version 5.1.6

Run on: December 4, 2003, 14:31:50 ; Search time 2.73874 Seconds
                 (without alignments)
                 741.553 Million cell updates/sec

Title: US-08-569-749-9
Perfect score: 295
Sequence: 1 PEQLASAGFYYVGRNDDVKC...............CWESGDDPWVEHAKWFPRCE 48

Scoring table:
Scoring table: BLOSUM62
                 Gapop 10.0 , Gapext 0.5

Searched: 328717 seqs, 42310858 residues

Total number of hits satisfying chosen parameters: 328717

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   Issued Patents AA:*
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                 2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
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                 4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
                 5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
                 6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Match DB	Length	DB	ID	Description
1	295	100.0	48	3	US-08-569-749-9	Sequence 9, Appli
2	295	100.0	48	5	PCT-US96-12860-9	Sequence 9, Appli
3	295	100.0	438	5	PCT-US95-05922A-2	Sequence 2, Appli
4	295	100.0	618	5	US-08-569-749-2	Sequence 2, Appli
5	295	100.0	618	4	US-09-069-023-29	Sequence 29, Appli
6	295	100.0	618	5	PCT-US96-12860-2	Sequence 2, Appli
7	283	95.9	68	2	US-08-511-485-27	Sequence 27, Appl
8	283	95.9	68	5	US-09-201-936-27	Sequence 27, Appl
9	283	95.9	618	2	US-08-511-485-8	Sequence 8, Appli
10	283	95.9	618	3	US-09-212-971-8	Sequence 8, Appli
11	283	95.9	618	3	US-08-800-929A-8	Sequence 8, Appli
12	283	95.9	283	4	US-09-617-053A-8	Sequence 8, Appli
13	283	95.9	618	4	US-09-201-936-8	Sequence 8, Appli
14	282	95.6	618	5	US-08-569-749-10	Sequence 10, Appl
15	282	95.6	48	5	PCT-US96-12860-10	Sequence 10, Appl
16	282	95.6	68	2	US-08-511-485-26	Sequence 26, Appl
17	282	95.6	68	4	US-09-201-936-26	Sequence 26, Appl
18	282	95.6	604	2	US-08-511-485-6	Sequence 6, Appli
19	282	95.6	604	3	US-09-212-971-6	Sequence 6, Appli
20	282	95.6	604	3	US-08-800-929A-6	Sequence 6, Appli
21	282	95.6	604	4	US-08-569-749-4	Sequence 4, Appli
22	282	95.6	604	4	US-09-617-053A-6	Sequence 6, Appli
23	282	95.6	604	4	US-09-201-936-6	Sequence 6, Appli
24	282	95.6	604	4	PCT-US96-12860-4	Sequence 4, Appl
25	282	95.6	612	3	US-09-212-971-14	Sequence 14, Appl
26	282	95.6	612	3	US-08-800-929A-14	Sequence 14, Appl
27	282	95.6	612	3	US-08-569-749-14	Sequence 14, Appl
28	282	95.6	612	4	US-09-617-053A-14	Sequence 14, Appl
29	282	95.6	612	5	PCT-US96-12860-14	Sequence 14, Appl
30	276	93.6	591	4	US-09-201-936-42	Sequence 42, Appl
31	268	90.8	600	3	US-09-212-971-12	Sequence 12, Appl
32	268	90.8	600	4	US-08-800-929A-12	Sequence 12, Appl
33	262	90.8	602	4	US-09-617-053A-12	Sequence 12, Appl
34	262	88.8	602	4	US-09-201-936-40	Sequence 40, Appl
35	231	78.3	50	3	US-08-975-080-31	Sequence 31, Appl
36	230	78.0	50	3	US-08-975-080-22	Sequence 22, Appl
37	230	78.0	50	4	US-08-975-080-30	Sequence 30, Appl
38	198	67.1	68	2	US-08-657-759-6	Sequence 6, Appli
39	198	67.1	68	4	US-09-511-485-28	Sequence 28, Appl
40	198	67.1	268	4	US-09-201-936-31	Sequence 31, Appl
41	198	67.1	268	3	US-08-836-134-22	Sequence 22, Appl
42	198	67.1	268	4	US-09-493-784-22	Sequence 22, Appl
43	195	66.1	298	4	US-09-127-928-2	Sequence 2, Appli
44	188	63.7	377	4	US-09-502-528-3	Sequence 3, Appli
45	187	63.4	50	3	US-08-975-080-21	Sequence 21, Appl

ALIGNMENTS

RESULT 1
US-08-569-749-9
Sequence 9, Application US/08569749
; Patent No. 6187557
; GENERAL INFORMATION:
; APPLICANT: Rothe, Mike
; APPLICANT: Goeddel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749
; FILING DATE:
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Brezner, David J.
; REGISTRATION NUMBER: 24,774
; REFERENCE/DOCKET NUMBER: A-62464/DJB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415)781-1989
; TELEFAX: (415)398-3249
; INFORMATION FOR SEQ ID NO: 9:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 48 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-569-749-9

Query Match   100.0%; Score 295; DB 3; Length 48;
Best Local Similarity   100.0%; Pred. No. 4.8e-30;
Matches   48; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
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Db   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48

RESULT 2
PCT-US96-12860-9
Sequence 9, Application PCT/US9612860
GENERAL INFORMATION:
   APPLICANT: TULARIK, INC.
   TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
   NUMBER OF SEQUENCES: 14
   CORRESPONDENCE ADDRESS:
      ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
      STREET: 4 Embarcadero Center, Suite 3400
      CITY: San Francisco
      STATE: California
      COUNTRY: USA
      ZIP: 94111
   COMPUTER READABLE FORM:
      MEDIUM TYPE: Floppy disk
      COMPUTER: IBM PC compatible
      OPERATING SYSTEM: PC-DOS/MS-DOS
      SOFTWARE: Patentin Release #1.0, Version #1.30
   CURRENT APPLICATION DATA:
      APPLICATION NUMBER: PCT/US96/12860
      FILING DATE: 06 AUG 1996
      CLASSIFICATION:
   PRIOR APPLICATION DATA:
      APPLICATION NUMBER: U.S. Serial Nos. 08/512,916 & 08/569,749
      CLASSIFICATION:
   ATTORNEY/AGENT INFORMATION:
      NAME: Brezner, David J.
      REGISTRATION NUMBER: 24,774
      REFERENCE/DOCKET NUMBER: A-62464/DJB
   TELECOMMUNICATION INFORMATION:
      TELEPHONE: (415)781-1989
      TELEFAX: (415)398-3249
INFORMATION FOR SEQ ID NO: 9:
   SEQUENCE CHARACTERISTICS:
      LENGTH: 48 amino acids
      TYPE: amino acid
      STRANDEDNESS: single
      TOPOLOGY: linear
   MOLECULE TYPE: protein
PCT-US96-12860-9

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Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
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Db   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
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Query Match   100.0%; Score 295; DB 5; Length 48;
Best Local Similarity 100.0%; Pred. No. 4.8e-30;
Matches   48; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

RESULT 3
PCT-US95-05922A-2
Sequence 2, Application PC/TUS505922A
GENERAL INFORMATION:
   APPLICANT: HE, ET AL.
   TITLE OF INVENTION: Human Inhibitor of Apoptosis Gene 1
   NUMBER OF SEQUENCES: 8
   CORRESPONDENCE ADDRESS:
      ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
                 CECCHI, STEWART & OLSTEIN
      STREET: 6 BECKER FARM ROAD
      CITY: ROSELAND
      STATE: NEW JERSEY
      COUNTRY: USA
      ZIP: 07068
   COMPUTER READABLE FORM:
      MEDIUM TYPE: 3.5 INCH DISKETTE
      COMPUTER: IBM PS/2
      OPERATING SYSTEM: MS-DOS
      SOFTWARE: WORD PERFECT 5.1
   CURRENT APPLICATION DATA:
      APPLICATION NUMBER: PCT/US95/05922A
      FILING DATE: 11 MAY 1995
      CLASSIFICATION:
   PRIOR APPLICATION DATA:
      APPLICATION NUMBER:
      FILING DATE:
   ATTORNEY/AGENT INFORMATION:
      NAME: Ferraro, Gregory D.
      REGISTRATION NUMBER: 36,134
      REFERENCE/DOCKET NUMBER: 325800-292
   TELECOMMUNICATION INFORMATION:
      TELEPHONE: 201-994-1744
      TELEFAX: 201-994-1700
INFORMATION FOR SEQ ID NO: 2:
   SEQUENCE CHARACTERISTICS:
      LENGTH: 438 AMINO ACIDS
      TYPE: AMINO ACID
      STRANDEDNESS:
      TOPOLOGY: LINEAR
   MOLECULE TYPE: PROTEIN
PCT-US95-05922A-2

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Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
        ||||||||||||||||||||||||||||||||||||||||||||||||
Db 107 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 154
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Query Match   100.0%; Score 295; DB 5; Length 438;
Best Local Similarity 100.0%; Pred. No. 5.3e-29;
Matches   48; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

RESULT 4
US-08-569-749-2
Sequence 2, Application US/08569749
Patent No. 6187557
GENERAL INFORMATION:
   APPLICANT: Rothe, Mike
   APPLICANT: Goeddel, David V
   TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
   NUMBER OF SEQUENCES: 14
   CORRESPONDENCE ADDRESS:
      ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
      STREET: 4 Embarcadero Center, Suite 3400
      CITY: San Francisco
      STATE: California
      COUNTRY: USA
      ZIP: 94111
   COMPUTER READABLE FORM:
      MEDIUM TYPE: Floppy disk
      COMPUTER: IBM PC compatible
      OPERATING SYSTEM: PC-DOS/MS-DOS
      SOFTWARE: Patentin Release #1.0, Version #1.30
   CURRENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/569,749
      FILING DATE:
      CLASSIFICATION: 514
   ATTORNEY/AGENT INFORMATION:
      NAME: Brezner, David J.
      REGISTRATION NUMBER: 24,774
      REFERENCE/DOCKET NUMBER: A-62464/DJB
   TELECOMMUNICATION INFORMATION:
      TELEPHONE: (415)781-1989
      TELEFAX: (415)398-3249
INFORMATION FOR SEQ ID NO: 2:
   SEQUENCE CHARACTERISTICS:
      LENGTH: 618 amino acids
      TYPE: amino acid
      STRANDEDNESS: single
      TOPOLOGY: linear
   MOLECULE TYPE: protein
US-08-569-749-2

Query Match   100.0%; Score 295; DB 3; Length 618;
Best Local Similarity 100.0%; Pred. No. 7.6e-29;

Qy   Matches   48; Conservative   0;   Mismatches   0;   Indels   0;   Gaps   0;

Db   287 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE   334

RESULT 5
US-09-069-023-29
; Sequence 29, Application US/09069023A
; Patent No. 6348573
; GENERAL INFORMATION:
; APPLICANT: Nunez, Gabriel
; APPLICANT: Inohara, Naohiro
; APPLICANT: Koseki, Takeyoshi
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR IDENTIFYING APOPTOSIS
;   TITLE OF INVENTION: SIGNALING PATHWAY INHIBITORS AND ACTIVATORS
; FILE REFERENCE: UM-03333
; CURRENT APPLICATION NUMBER: US/09/069,023A
; CURRENT FILING DATE: 1998-04-27
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Patentin Ver. 2.0
;
; SEQ ID NO 29
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens

Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE   48
         ||||||||||||||||||||||||||||||||||||||||||||||||
Db   287 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE   334

Query Match   100.0%; Score 295; DB 4;   Length 618;
Best Local Similarity   100.0%; Pred. No. 7.6e-29;
Matches   48; Conservative   0;   Mismatches   0;   Indels   0;   Gaps   0;

RESULT 6
PCT-US96-12860-2
; Sequence 2, Application PC/TUS9612860
; GENERAL INFORMATION:
; APPLICANT: TULARIK, INC.
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US96/12860
; FILING DATE: 06 AUG 1996
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: U.S. Serial Nos. 08/512,946 & 08/569,749
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Brezner, David J.
; REGISTRATION NUMBER: 24,774
; REFERENCE/DOCKET NUMBER: A-62464/DJB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415)781-1989
; TELEFAX: (415)398-3249
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 618 amino acids

; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein

Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE   48
         ||||||||||||||||||||||||||||||||||||||||||||||||
Db   287 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE   334

Query Match   100.0%; Score 295; DB 5;   Length 618;
Best Local Similarity   100.0%; Pred. No. 7.6e-29;
Matches   48; Conservative   0;   Mismatches   0;   Indels   0;   Gaps   0;

RESULT 7
US-08-511-485-27
; Sequence 27, Application US/08511485
; Patent No. 5919912
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mackenzie, Alexander E.
; APPLICANT: Baird, Stephen
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; NUMBER OF SEQUENCES: 38
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson P.C.
; STREET: 225 Franklin Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110-2804
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/511,485
; FILING DATE: 04-AUG-1995
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Clark, Paul T.
; REGISTRATION NUMBER: 30,162
; REFERENCE/DOCKET NUMBER: 07540/002001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617/542-5070
; TELEFAX: 617/542-8906
; TELEX: 200154
; INFORMATION FOR SEQ ID NO: 27:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 68 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: both
; MOLECULE TYPE: protein

Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE   48
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Db   287 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE   334

Query Match   100.0%; Score 295; DB 2;   Length 68;
Best Local Similarity   100.0%; Pred. No. 2.2e-28;
Matches   48; Conservative   0;   Mismatches   1;   Indels   0;   Gaps   0;

RESULT 8
US-09-201-936-27
; Sequence 27, Application US/09201936
; Patent No. 6541457
; GENERAL INFORMATION:

Qy   1 PEQLASAGFYYVGRNDDVKCFCPGCDGGLRCWESGDDPWVEHAKWFPRCE   48
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Db   19 PEQLASAGFYYVGRNDDVKCFCPGCDGGLRCWESGDDPWVEHAKWFPRCE   66

Query Match   95.9%; Score 283; DB 2;   Length 68;
Best Local Similarity   97.9%; Pred. No. 2.0e-28;
Matches   47; Conservative   0;   Mismatches   1;   Indels   0;   Gaps   0;

APPLICANT: Korneluk, Robert G.
APPLICANT: MacKenzie, Alexander E.
APPLICANT: Baird, Stephen
APPLICANT: Liston, Peter
TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
PROBES, AND DETECTION METHODS
FILE REFERENCE: 07891/003003
CURRENT APPLICATION NUMBER: US/09/201,936
CURRENT FILING DATE: 1998-12-01
EARLIER APPLICATION NUMBER: 09/011,356
EARLIER FILING DATE: 1998-02-04
EARLIER APPLICATION NUMBER: 08/511,485
EARLIER FILING DATE: 1995-08-04
EARLIER APPLICATION NUMBER: PCT/IB96/01022
EARLIER FILING DATE: 1996-08-05
EARLIER APPLICATION NUMBER: 08/576,956
EARLIER FILING DATE: 1995-12-22
SOFTWARE: FastSEQ for Windows Version 3.0
NUMBER OF SEQ ID NOS: 45
SEQ ID NO 27
LENGTH: 68
TYPE: PRT
ORGANISM: Homo sapiens
US-09-201-936-27

Query Match   95.9%;   Score 283;   DB 4;   Length 68;
Best Local Similarity   97.9%;   Pred. No. 2.2e-28;
Matches 47;   Conservative 0;   Mismatches 1;   Indels 0;   Gaps 0;

Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
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Db   19 PEQLASAGFYYVGRNDDVKCFGCDGGLRCWESGDDPWVEHAKWFPRCE 66

RESULT 9
US-08-511-485-8
; Sequence 8, Application US/08511485
; Patent No. 5919912
GENERAL INFORMATION:
APPLICANT: Korneluk, Robert G.
APPLICANT: MacKenzie, Alexander E.
APPLICANT: Baird, Stephen
TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
PROBES, AND DETECTION METHODS
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/511,485
FILING DATE: 04-AUG-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 07540/002001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 618 amino acids
TYPE: amino acid

STRANDEDNESS: not relevant
TOPOLOGY: both
MOLECULE TYPE: protein
US-08-511-485-8

Query Match   95.9%;   Score 283;   DB 2;   Length 618;
Best Local Similarity   97.9%;   Pred. No. 2.4e-27;
Matches 47;   Conservative 0;   Mismatches 1;   Indels 0;   Gaps 0;

Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
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Db 287 PEQLASAGFYYVGRNDDVKCFGCDGGLRCWESGDDPWVEHAKWFPRCE 334

RESULT 10
US-09-212-971-8
; Sequence 8, Application US/09212971B
; Patent No. 6107041
GENERAL INFORMATION:
APPLICANT: Korneluk, Robert G
APPLICANT: MacKenzie, Alexander E
APPLICANT: Liston, Peter
APPLICANT: Baird, Stephen
APPLICANT: Tsang, Benjamin K
APPLICANT: Pratt, Christine
TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
DISEASE
FILE REFERENCE: 07891/009002
CURRENT APPLICATION NUMBER: US/09/212,971B
CURRENT FILING DATE: 1998-12-16
EARLIER APPLICATION NUMBER: 60/017,354
EARLIER FILING DATE: 1996-04-26
EARLIER APPLICATION NUMBER: 60/030,590
EARLIER FILING DATE: 1996-11-14
EARLIER APPLICATION NUMBER: 08/800,929
EARLIER FILING DATE: 1997-02-13
NUMBER OF SEQ ID NOS: 17
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
LENGTH: 618
TYPE: PRT
ORGANISM: Homo sapiens
US-09-212-971-8

RESULT 11
US-08-800-929A-8
; Sequence 8, Application US/08800929A
; Patent No. 6133437
GENERAL INFORMATION:
APPLICANT: Korneluk, Robert G
APPLICANT: MacKenzie, Alexander E
APPLICANT: Liston, Peter
APPLICANT: Baird, Stephen
APPLICANT: Tsang, Benjamin K
APPLICANT: Pratt, Christine
TITLE OF INVENTION: DETECTION AND MODULATION OF
IAPS AND NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
DISEASE
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Street
CITY: Boston

Query Match   95.9%;   Score 283;   DB 3;   Length 618;
Best Local Similarity   97.9%;   Pred. No. 2.4e-27;
Matches 47;   Conservative 0;   Mismatches 1;   Indels 0;   Gaps 0;

Qy   1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
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Db 287 PEQLASAGFYYVGRNDDVKCFGCDGGLRCWESGDDPWVEHAKWFPRCE 334

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;         STATE: MA
;       COUNTRY: USA
;           ZIP: 02110
; COMPUTER READABLE FORM:
;   MEDIUM TYPE: Diskette
;      COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
;      SOFTWARE: FastSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/800,929A
;  FILING DATE: 13-FEB-1997
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/030,590
;  FILING DATE: 14-NOV-1996
; APPLICATION NUMBER: 60/017,354
;  FILING DATE: 26-APR-1996
; ATTORNEY/AGENT INFORMATION:
;          NAME: Bieker-Brady, Kristina
; REGISTRATION NUMBER:
; REFERENCE/DOCKET NUMBER: 07891/009001
; TELECOMMUNICATION INFORMATION:
;     TELEPHONE: 617-428-0200
;      TELEFAX: 617-428-7045
;        TELEX:
; INFORMATION FOR SEQ ID NO:  8:
; SEQUENCE CHARACTERISTICS:
;        LENGTH: 618 amino acids
;          TYPE: amino acid
;  STRANDEDNESS: single
;      TOPOLOGY: linear
; MOLECULE TYPE: protein

US-08-800-929A-8

Qy          1 PEQLASAGFYVVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE  48
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Db        287 PEQLASAGFYVVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 334
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Query Match             95.9%;  Score 283;  DB 3;  Length 618;
Best Local Similarity   97.9%;  Pred. No. 2.4e-27;
Matches   47;  Conservative  0;  Mismatches  1;  Indels  0;  Gaps  0;

RESULT 12
US-09-617-053A-8
; Sequence 8, Application US/0961705JA
; Patent No. 6300492
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander B
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009003
; CURRENT APPLICATION NUMBER: US/09/617,053A
; CURRENT FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 08/800,929
; PRIOR FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 8
;        LENGTH: 618
;          TYPE: PRT
;      ORGANISM: Homo sapiens
;

US-09-617-053A-8

Query Match             95.9%;  Score 283;  DB 4;  Length 618;
Best Local Similarity   97.9%;  Pred. No. 2.4e-27;
Matches   47;  Conservative  0;  Mismatches  1;  Indels  0;  Gaps  0;
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Qy          1 PEQLASAGFYVVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE  48
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Db        287 PEQLASAGFYVVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 334
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RESULT 13
US-09-201-936-8
; Sequence 8, Application US/09201936
; Patent No. 6541457
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: MacKenzie, Alexander B.
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; FILE REFERENCE: 07891/003003
; CURRENT APPLICATION NUMBER: US/09/201,936
; CURRENT FILING DATE: 1998-12-01
; EARLIER APPLICATION NUMBER: 09/011,356
; EARLIER FILING DATE: 1998-02-04
; EARLIER APPLICATION NUMBER: PCT/IB96/01022
; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 08/576,956
; EARLIER FILING DATE: 1995-12-22
; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 8
;        LENGTH: 618
;          TYPE: PRT
;      ORGANISM: Homo sapiens

US-09-201-936-8

Query Match             95.9%;  Score 283;  DB 4;  Length 618;
Best Local Similarity   97.9%;  Pred. No. 2.4e-27;
Matches   47;  Conservative  0;  Mismatches  1;  Indels  0;  Gaps  0;
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Qy          1 PEQLASAGFYVVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE  48
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Db        287 PEQLASAGFYVVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 334
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RESULT 14
US-08-569-749-10
; Sequence 10, Application US/08569749
; Patent No. 6187557
; GENERAL INFORMATION:
; APPLICANT: Rothe, Mike
; APPLICANT: Goeddel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749
; FILING DATE:
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Brezner, David J.
; REGISTRATION NUMBER: 24,774
```

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;       REFERENCE/DOCKET NUMBER: A-62464/DJB
;       TELECOMMUNICATION INFORMATION:
;          TELEPHONE:    (415)781-1989
;          TELEFAX:      (415)398-3249
;    INFORMATION FOR SEQ ID NO: 10:
;       SEQUENCE CHARACTERISTICS:
;          LENGTH: 48 amino acids
;          TYPE: amino acid
;          STRANDEDNESS: single
;          TOPOLOGY: linear
;       MOLECULE TYPE: protein
US-08-569-749-10

Qy    Query Match         95.6%;   Score 282;  DB 3;   Length 48;
      Best Local Similarity   93.8%;   Pred. No. 2e-28;
      Matches    45;   Conservative    2;   Mismatches    1;   Indels   0;   Gaps   0;

Qy  1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
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Db  1 PEQLASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRCE 48

RESULT 15
PCT-US96-12860-10
;  Sequence 10, Application PC/TUS9612860
;  GENERAL INFORMATION:
;     APPLICANT: TULARIK, INC.
;     TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
;     NUMBER OF SEQUENCES: 14
;     CORRESPONDENCE ADDRESS:
;        ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
;        STREET: 4 Embarcadero Center, Suite 3400
;        CITY: San Francisco
;        STATE: California
;        COUNTRY: USA
;        ZIP: 94111
;     COMPUTER READABLE FORM:
;        MEDIUM TYPE: Floppy disk
;        COMPUTER: IBM PC compatible
;        OPERATING SYSTEM: PC-DOS/MS-DOS
;        SOFTWARE: PatentIn Release #1.0, Version #1.30
;     CURRENT APPLICATION DATA:
;        APPLICATION NUMBER: PCT/US96/12860
;        FILING DATE: 06 AUG 1996
;     PRIOR APPLICATION DATA:
;        APPLICATION NUMBER: U.S. Serial Nos. 08/512,946 & 08/569,749
;        CLASSIFICATION:
;     ATTORNEY/AGENT INFORMATION:
;        NAME: Brezner, David J.
;        REGISTRATION NUMBER: 24,774
;        REFERENCE/DOCKET NUMBER: A-62464/DJB
;     TELECOMMUNICATION INFORMATION:
;        TELEPHONE:    (415)781-1989
;        TELEFAX:      (415)398-3249
;  INFORMATION FOR SEQ ID NO: 10:
;     SEQUENCE CHARACTERISTICS:
;        LENGTH: 48 amino acids
;        TYPE: amino acid
;        STRANDEDNESS: single
;        TOPOLOGY: linear
;     MOLECULE TYPE: protein
PCT-US96-12860-10

Qy    Query Match         95.6%;   Score 282;  DB 5;   Length 48;
      Best Local Similarity   93.8%;   Pred. No. 2e-28;
      Matches    45;   Conservative    2;   Mismatches    1;   Indels   0;   Gaps   0;

Qy  1 PEQLASAGFYYVGRNDDVKCFCCDGGLRCWESGDDPWVEHAKWFPRCE 48
        |||||||||||||||:||||||||||||||||||||||||||||||||
Db  1 PEQLASAGFYYVGNSDDVKCFCCDGGLRCWESGDDPWVQHAKWFPRCE 48
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GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: December 4, 2003, 12:42:59 ; Search time 165 seconds
(without alignments)
6925.703 Million cell updates/sec

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :
Issued_Patents NA:*
1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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Title: US-08-569-749-1
Perfect Score: 2589
Sequence: 1 TCTAAGTAGTATCTTGGAAAA.........AAAAAAAAAAAAAAAAAAA 2589
Scoring table: IDENTITY NUC
Gapop 10.0 , Gapext 1.0
Searched: 569978 seqs, 220691566 residues

Total number of hits satisfying chosen parameters: 1139956

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
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2	2589	100.0	2589	5	PCT-US96-12860-1	Sequence 1, Appli
3	2535.8	97.9	3532	2	US-09-205-204-1	Sequence 1, Appli
4	2535.8	97.9	3732	3	US-09-212-971-7	Sequence 7, Appli
5	2535.8	97.9	3732	4	US-08-800-929A-7	Sequence 7, Appli
6	2535.8	97.9	3732	4	US-09-617-053A-7	Sequence 7, Appli
7	2514.8	97.1	2580	2	US-08-511-485-7	Sequence 7, Appli
8	2514.8	97.1	2580	4	US-09-201-936-7	Sequence 7, Appli
9	1430.2	55.2	1435	3	PCT-US95-05922A-1	Sequence 1, Appli
10	1345.2	52.0	2862	3	US-08-569-749-13	Sequence 13, Appli
11	1345.2	52.0	2862	5	PCT-US96-12860-13	Sequence 13, Appli
12	1343.6	51.9	3151	3	US-09-212-971-13	Sequence 13, Appli
13	1343.6	51.9	3151	4	US-08-800-929A-13	Sequence 13, Appl
14	1343.6	51.9	3151	4	US-09-617-053A-13	Sequence 13, Appl
15	1330.4	51.4	2416	4	US-09-201-936-41	Sequence 41, Appl
16	1157	44.7	3076	2	US-09-205-144-1	Sequence 1, Appl
17	1153.6	44.6	2563	4	US-09-016-434-1076	Sequence 1076, Ap
18	1153.6	44.6	2601	3	US-08-569-749-3	Sequence 3, Appli
19	1153.6	44.6	2601	5	PCT-US96-12860-3	Sequence 3, Appli
20	1152.2	44.5	6669	3	US-09-212-971-5	Sequence 5, Appli
21	1152.2	44.5	6669	4	US-08-800-929A-5	Sequence 5, Appli
22	1152.2	44.5	6669	4	US-09-617-053A-5	Sequence 5, Appli
23	1149	44.4	2676	2	US-08-511-485-5	Sequence 5, Appli
24	1149	44.4	2676	4	US-09-201-936-5	Sequence 5, Appli
25	931.6	36.0	2676	3	US-09-212-971-11	Sequence 11, Appl
26	931.6	36.0	2676	4	US-08-800-929A-11	Sequence 11, Appl
27	931.6	36.0	2676	4	US-09-617-053A-11	Sequence 11, Appl
28	921.8	35.6	2450	4	US-09-201-936-39	Sequence 39, Appl
29	137.4	5.3	2691	3	US-09-212-971-9	Sequence 9, Appli
30	137.4	5.3	2691	3	US-08-800-929A-9	Sequence 9, Appli
31	137.4	5.3	2691	4	US-09-617-053A-9	Sequence 9, Appli
32	136.8	5.3	1588	4	US-09-016-434-1053	Sequence 1053, Ap
33	136.8	5.3	1659	3	US-09-239-867-3	Sequence 3, Appli
34	136.8	5.3	5232	3	US-09-212-971-3	Sequence 3, Appli
35	136.8	5.3	5232	4	US-08-800-929A-3	Sequence 3, Appli
36	136.8	5.3	5232	4	US-09-617-053A-3	Sequence 3, Appli
37	136.6	5.3	2100	4	US-08-511-485-9	Sequence 9, Appli
38	136.6	5.3	2100	4	US-09-201-936-9	Sequence 9, Appli
39	135.2	5.2	2540	2	US-08-511-485-3	Sequence 3, Appli
40	135.2	5.2	2540	4	US-09-392-580-1	Sequence 1, Appli
41	135.2	5.2	2540	4	US-09-201-936-3	Sequence 3, Appli
42	92.8	3.6	1559	4	US-09-239-867-1	Sequence 1, Appli
43	80.8	3.1	1337	4	US-09-127-928-1	Sequence 1, Appli
44	76.4	3.0	1134	4	US-09-502-528-2	Sequence 2, Appli
45	76.4	3.0	1739	4	US-09-502-528-1	Sequence 1, Appli

ALIGNMENTS

RESULT 1
US-08-569-749-1

; Sequence 1, Application US/08569749
; Patent No. 6187557
; GENERAL INFORMATION:
; APPLICANT: Rothe, Mike
; APPLICANT: Goeddel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749
; FILING DATE:
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Brezner, David J.
; REGISTRATION NUMBER: 24,774
; REFERENCE/DOCKET NUMBER: A-62464/DJB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415)781-1989
; TELEFAX: (415)398-3249
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2589 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-569-749-1

Query Match 100.0%; Score 2589; DB 3; Length 2589;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 2589; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TCTAAGTAGTATCTTGGAAAATTCAGAGAGATTACTCATCCTTACCTGAATATAAAACTGAGAT 60
      ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1 TCTAAGTAGTATCTTGGAAAATTCAGAGAGATTACTCATCCTTACCTGAATATAAAACTGAGAT 60

Qy 61 AAATCCAGTAAAAGAAAAGTGTAGTAAATTCTRACATAAGAGTCTRTCATTGATTTCTTTTGG 120

```
Qy        61 AAAATCCAGTAAAGAAAAGTGTAGTAAAATCTTACATAAAGAGTCTATCATTGATTTCTTTGG 120
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        61 AAAATCCAGTAAAGAAAAGTGTAGTAAAATCTTACATAAAGAGTCTATCATTGATTTCTTTGG 120

Qy       121 TGGTAAAAATCTTAGTTCATGTGAAGAAATTCATGTGAATGTTTTAGCTATCAAACAGC 180
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       121 TGGTAAAAATCTTAGTTCATGTGAAGAAATTCATGTGAATGTTTTAGCTATCAAACAGC 180

Qy       181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTCGTATC 240
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTCGTATC 240

Qy       241 AAAACATTAAGAGTATAATGGAAGATAGCACGGATCTTGTCAGATTGGACAAAACAGCAACA 300
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       241 AAAACATTAAGAGTATAATGGAAGATAGCACGGATCTTGTCAGATTGGACAAAACAGCAACA 300

Qy       301 AACAAAAAAATGAAGTAATGACTTTTTCCTTGTGAACTCTTACAGAAATGTTCTACATATTCAACTT 360
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       301 AACAAAAAAATGAAGTAATGACTTTTTCCTTGTGAACTCTTACAGAAATGTTCTACATATTCAACTT 360

Qy       361 TCCCCGCCCGGGGGTGCCTGGCCTTGCTTCAGAAAGGAGTCTTGCTCGTGCTGGTTTTTTATTATACTG 420
             |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       361 TCCCCGCCCGGGGGTGCCTGGCCTTGCTTCAGAAAGGAGTCTTGCTCGTGCTGGTTTTTTATTATACTG 420

Qy       421 GTGTGAATGACAAGGTCAAATGCTTCTGTTGTTGGTGGCCTGGATAACTGGAAAACTTAG 480
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       421 GTGTGAATGACAAGGTCAAATGCTTCTGTTGTTGGTGGCCTGGATAACTGGAAAACTTAG 480

Qy       481 GAGACAGTGCCTATTCAAAAAGCATAAACAGCTATATCCTAGCTGTGTAGCTTTATTCAGAAATC 540
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       481 GAGACAGTGCCTATTCAAAAAGCATAAACAGCTATATCCTAGCTGTGTAGCTTTATTCAGAAATC 540

Qy       541 TGGTTTCAGCTAGTCTGGGGATCCAGCCACCTCAACTTCCAAATGAGAAACAGTTTTTG 600
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       541 TGGTTTCAGCTAGTCTGGGGATCCAGCCACCTCAACTTCCAAATGAGAAACAGTTTTTG 600

Qy       601 CACATTCATTATCTCCCACCTTGGGAACATAGTAGCTTGTTCAGTGGTTCTTACTTCCAGCC 660
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       601 CACATTCATTATCTCCCACCTTGGGAACATAGTAGCTTGTTCAGTGGTTCTTACTTCCAGCC 660

Qy       661 TTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACTCTTCTCATCGGAGGACTAACC 720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       661 TTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACTCTTCTCATCGGAGGACTAACC 720

Qy       721 CCTACAGTTATGCAATGAGTACTGAAGAGCCAGATTTCTTACCTACCATATGTGGCCAT 780
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       721 CCTACAGTTATGCAATGAGTACTGAAGAGCCAGATTTCTTACCTACCATATGTGGCCAT 780

Qy       781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAAGCTGGTGTTTTATTATATAGGACCTGGGAG 840
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAAGCTGGTGTTTTATTATATAGGACCTGGGAG 840

Qy       841 ATAGGGTAGCCTGCTTTGCCTGTGGTGGGAAGCTCAGTACTTGAGTAACCCAAAGGATGATG 900
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       841 ATAGGGTAGCCTGCTTTGCCTGTGGTGGGAAGCTCAGTACTTGAGTAACCCAAAGGATGATG 900

Qy       901 CTATGTCAGAACACCGGAGGCATTTCCCAACTGTCCAATTTGGAAAATTCTCTAGAAA 960
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       901 CTATGTCAGAACACCGGAGGCATTTCCCAACTGTCCAATTTGGAAAATTCTCTAGAAA 960

Qy       961 CTCTGAGGTTTAGCATTTCAAATCTGACGATGCAGCAGCTTCGAGTGCATGCAATGAGAACAT 1020
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       961 CTCTGAGGTTTAGCATTTCAAATCTGACGATGCAGCAGCTTCGAGTGCATGCAATGAGAACAT 1020

Qy      1021 TTATGTACTGGGCCATCTAGTGTTCCAGCTGAGCACCATGACCTTGCAAGTGCTGGTTTTT 1080
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1021 TTATGTACTGGGCCATCTAGTGTTCCAGCTGAGCACCATGACCTTGCAAGTGCTGGTTTTT 1080

Qy      1081 ATTATGTGGGGTCGCAAATGATGATGCTTTTGTTTGTTGTGATGGTGGGCTTGAGGTGTT 1140
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1081 ATTATGTGGGGTCGCAAATGATGATGCTTTTGTTTGTTGTGATGGTGGGCTTGAGGTGTT 1140

Qy      1141 GGGAATCTGAGATGATCCATGGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 1200
             |||||||| || | ||
Db      1141 GGGAATCTGAGATGATCCATGGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 1200
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Db      1141 GGGAATCTGAGATGATCCATGGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 1200
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy      1201 TGATACGGAATGAAAAGGCCAAGAGAGTTTGTTGATGAGATTCAAGGTAGATATCCTCATCTTC 1260
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1201 TGATACGGAATGAAAAGGCCAAGAGAGTTTGTTGATGAGATTCAAGGTAGATATCCTCATCTTC 1260

Qy      1261 TTGAACAGCTGTTGTCAACTTCAGATACCACTGGAGAACGAAAAATGCTGACCCACCAATTA 1320
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1261 TTGAACAGCTGTTGTCAACTTCAGATACCACTGGAGAACGAAAAATGCTGACCCACCAATTA 1320

Qy      1321 TTCATTTTGGACCTGGAGAAGATTTCTTCAGAAGATGCTGTCATGATGAACTACACCTGTGG 1380
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1321 TTCATTTTGGACCTGGAGAAGATTTCTTCAGAAGATGCTGTCATGATGAACTACACCTGTGG 1380

Qy      1381 TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGGACCTGGTGAAACAGTTCAAAGTA 1440
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1381 TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGGACCTGGTGAAACAGTTCAAAGTA 1440

Qy      1441 AAATCCTGACAACTGGGAGGAACTATAAAAACAGTTAAATGAATATTGTGTCAGCACTTCTAA 1500
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1441 AAATCCTGACAACTGGGAGGAACTATAAAAACAGTTAAATGAATATTGTGTCAGCACTTCTAA 1500

Qy      1501 ATGCTGAAGATGAAGATGAAAAAGAGAGGAGGAGGAGGAGAAAAAAACAAGCTGAAGAAAATGGCATCAG 1560
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1501 ATGCTGAAGATGAAGATGAAAAAGAGAGGAGGAGGAGGAGAAAAAAACAAGCTGAAGAAAATGGCATCAG 1560

Qy      1561 ATGATTTTGTCATTAAATTCGGAGAACAGAAATGGCTCTTCTTCAACAATTGACATTGGTGTGC 1620
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1561 ATGATTTTGTCATTAAATTCGGAGAACAGAAATGGCTCTTCTTCAACAATTGACATTGGTGTGC 1620

Qy      1621 TTCCTATCCTGGATAATCTTTTAAAGCCAAATGTAATTAATTAATAAACAGGAACACGAGATAATTA 1680
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1621 TTCCTATCCTGGATAATCTTTTAAAGCCAAATGTAATTAATTAATAAACAGGAACACGAGATAATTA 1680

Qy      1681 TTAAACAAAAAACACAGATACCTTTACAAGCGAGAGAACTGATTGATACCATTTTGGTTA 1740
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1681 TTAAACAAAAAACACAGATACCTTTACAAGCGAGAGAACTGATTGATACCATTTTGGTTA 1740

Qy      1741 AAGGAAAATGCTGCGGGCCAACACATCTTCAAAAAACTGTCTAAAAGAAATTGACTCTACATTGT 1800
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1741 AAGGAAAATGCTGCGGGCCAACACATCTTCAAAAAACTGTCTAAAAGAAATTGACTCTACATTGT 1800

Qy      1801 ATAAGAACTTATTTGTGGATAAGAATATGAAGTAATATTCCAACAGAACAAGAAGTGTTTCAGGTC 1860
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1801 ATAAGAACTTATTTGTGGATAAGAATATGAAGTAATATTCCAACAGAACAAGAAGTGTTTCAGGTC 1860

Qy      1861 TGTCACTGGAAGAACAAATTGAGGAGGTTGCAAGAACGAACTTGTAAAGTGTGTATGG 1920
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1861 TGTCACTGGAAGAACAAATTGAGGAGGTTGCAAGAACGAACTTGTAAAGTGTGTATGG 1920

Qy      1921 ACAAAGAAGAGTTTCGTGTTGTATTTATTCCTTGTGTGGTCATCCTGGTAGTATGCCAGGAAATGTG 1980
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1921 ACAAAGAAGAGTTTCGTGTTGTATTTATTCCTTGTGTGGTCATCCTGGTAGTATGCCAGGAAATGTG 1980

Qy      1981 CCCCTTCTCTAAGAAAAATGCCCTATTTGCAGGGGGTATAATCAAGGGTACTGTTCGTACAT 2040
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Db      1981 CCCCTTCTCTAAGAAAAATGCCCTATTTGCAGGGGGTATAATCAAGGGTACTGTTCGTACAT 2040

Qy      2041 TTCTCTCTTAAAGAAAAAAAAGTCTATATTTTTAACCTGGCGATAAAAAAGGTCTTTAAAAATATT 2100
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2041 TTCTCTCTTAAAGAAAAAAAAGTCTATATTTTTAACCTGGCGATAAAAAAGGTCTTTAAAAATATT 2100

Qy      2101 GTTGAAACACTTGAAGGCCATCTAAAGTAAAAGAAATTATATGAGTTTTTCAAATTAGTAAACA 2160
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2101 GTTGAAACACTTGAAGGCCATCTAAAGTAAAAGAAATTATATGAGTTTTTCAAATTAGTAAACA 2160

Qy      2161 TTCATGTTCTAGTCTGCTTTGGTACTAATAATCTGTTTCTGAAAAGATAGGTATCATATA 2220
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2161 TTCATGTTCTAGTCTGCTTTGGTACTAATAATCTGTTTCTGAAAAGATAGGTATCATATA 2220

Qy      2221 TTTAATCTTAAATCTGTTTTATTTAACAAGGGAAGAGATTTATGTTTTGTGTGAACTATATATTAGTTAT 2280
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2221 TTTAATCTTAAATCTGTTTTATTTAACAAGGGAAGAGATTTATGTTTTGTGTGAACTATATATTAGTTAT 2280
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RESULT 2
PCT-US96-12860-1
Sequence 1, Application PC/TUS9612860
GENERAL INFORMATION:
  APPLICANT: TULARIK, INC.
  TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
  NUMBER OF SEQUENCES: 14
  CORRESPONDENCE ADDRESS:
  ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
  STREET: 4 Embarcadero Center, Suite 3400
  CITY: San Francisco
  STATE: California
  COUNTRY: USA
  ZIP: 94111
  COMPUTER READABLE FORM:
  MEDIUM TYPE: Floppy disk
  COMPUTER: IBM PC compatible
  OPERATING SYSTEM: PC-DOS/MS-DOS
  SOFTWARE: PatentIn Release #1.0, Version #1.30
  CURRENT APPLICATION DATA:
  APPLICATION NUMBER: PCT/US96/12860
  FILING DATE: 06 AUG 1996
  CLASSIFICATION:
  PRIOR APPLICATION DATA:
  APPLICATION NUMBER: U.S. Serial Nos. 08/512,945 & 08/569,749
  CLASSIFICATION:
  ATTORNEY/AGENT INFORMATION:
  NAME: Brezner, David J.
  REGISTRATION NUMBER: 24,774
  REFERENCE/DOCKET NUMBER: A-62464/DJB
  TELECOMMUNICATION INFORMATION:
  TELEPHONE: (415)781-1989
  TELEFAX: (415)398-3249
INFORMATION FOR SEQ ID NO: 1:
  SEQUENCE CHARACTERISTICS:
  LENGTH: 2589 base pairs
  TYPE: nucleic acid
  STRANDEDNESS: single
  TOPOLOGY: linear
  MOLECULE TYPE: cDNA
PCT-US96-12860-1

Query Match 100.0%;   Score 2589;   DB 5;   Length 2589;
Best Local Similarity 100.0%;   Pred. No. 0;
Matches 2589;   Mismatches 0;   Conservative 0;   Indels 0;   Gaps 0;

(Sequence alignment — Qy = query, Db = database; | denotes identity)

Qy   1 TCTAAGTAGTATCTTGGAAATTCAGAGAGATACTCATCCTACCTGAATATAAAACTGAGAT   60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1 TCTAAGTAGTATCTTGGAAATTCAGAGAGATACTCATCCTACCTGAATATAAAACTGAGAT   60

Qy   61 AAATCCAGTAAAAGAAAGTGTAGTAAAATTCTACATAAGAAGTCTATCATTGATTCTTTTGG   120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 AAATCCAGTAAAAGAAAGTGTAGTAAAATTCTACATAAGAAGTCTATCATTGATTCTTTTGG   120

Qy   121 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAATGTTTAGCTATTGAAACAGC   180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAATGTTTAGCTATTGAAACAGC   180

Qy   181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGCTCCCTCGTGTATC   240
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Db   181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGCTCCCTCGTGTATC   240

Qy   241 AAAACATTAAGAGTATAATGGAAGATAGCACGGATCTTGTCAGATTGGACAAAACAGCAACA   300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241 AAAACATTAAGAGTATAATGGAAGATAGCACGGATCTTGTCAGATTGGACAAAACAGCAACA   300

Qy   301 AACAAAAAATGAAGTAATGACTTTTCCTGTGAAACTCTACACAGAATTCTACATATTCAACTT   360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   301 AACAAAAAATGAAGTAATGACTTTTCCTGTGAAACTCTACACAGAATTCTACATATTCAACTT   360

Qy   361 TCCCCGCCGCGGGGTGCCCTGTCTCAGAAAAGGAGTCTTGCTCGTGCTGGTTTTTATTATACTG   420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   361 TCCCCGCCGCGGGGTGCCCTGTCTCAGAAAAGGAGTCTTGCTCGTGCTGGTTTTTATTATACTG   420

Qy   421 GTGTGAATGACAAGGTCAAATGCTTCTGTTGTGGCGGTGGATAACTGGAACTAG   480
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Qy   481 GAGACAGTCCTATTCAAAAGCATAAAACAGCTATATCCTAGCTGTAGCTTTATTCAGAATC   540
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Db   481 GAGACAGTCCTATTCAAAAGCATAAAACAGCTATATCCTAGCTGTAGCTTTATTCAGAATC   540

Qy   541 TGGTTTCAGCTAGTCTGGGGATCCACCTCTAAGAATAACAGTCTCCAAATGAGAAACAGTTTG   600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   541 TGGTTTCAGCTAGTCTGGGGATCCACCTCTAAGAATAACAGTCTCCAAATGAGAAACAGTTTG   600

Qy   601 CACATTCATTATCTCCCACCTTGGAACATAGTAGTTCAGTGGTTCATCGAGGACTAACC   660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   601 CACATTCATTATCTCCCACCTTGGAACATAGTAGTTCAGTGGTTCATCGAGGACTAACC   660

Qy   661 TTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACAGTCCATTAATTCCCGAGGACTAACC   720
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Db   661 TTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACAGTCCATTAATTCCCGAGGACTAACC   720

Qy   721 CCTACAGTTATGCAAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCATATGTGCCAT   780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   721 CCTACAGTTATGCAAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCATATGTGCCAT   780

Qy   781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATTAGGAGCCTGGAG   840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATTAGGAGCCTGGAG   840

Qy   841 ATAGGGTAGCCTGCTTTGCCTGTGGGGAGGCAAACCAAAGGATGATG   900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   841 ATAGGGTAGCCTGCTTTGCCTGTGGGGAGGCAAACCAAAGGATGATG   900

Qy   901 CTATGTCAGAACACCGGAGGCATTTCCCAACTGTCCAATTTTGGAAAATTCTCTAGAA   960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   901 CTATGTCAGAACACCGGAGGCATTTCCCAACTGTCCAATTTTGGAAAATTCTCTAGAA   960

Qy   961 CTCTGAGGTTTAGCATTTCAAAATCTGAGCAATGCAGAGACACATGCAGAAGCATCGAGAACAT 1020
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   961 CTCTGAGGTTTAGCATTTCAAAATCTGAGCAATGCAGAGACACATGCAGAAGCATCGAGAACAT 1020

Qy 1021 TTATGTACTGGCCATCTAGTGTTCAGCCTGAGCAGCTTGCCAAGTGCTCGGTTTTT 1080
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1021 TTATGTACTGGCCATCTAGTGTTCAGCCTGAGCAGCTTGCCAAGTGCTCGGTTTTT 1080

Qy 1081 ATTATGTGGGGTCGCGAATGATGTCAAATGCTTTTGTTGTGTATGGGTTGCTTGAGGTGTT 1140
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Db 1081 ATTATGTGGGGTCGCGAATGATGTCAAATGCTTTTGTTGTGTATGGGTTGCTTGAGGTGTT 1140

Qy 2281 GTATGTGTACCTAAGGGAGTAGTGAGTTCACTGCTTGTTATGTATCATTTCAGGAGTTACTGG 2340
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 2281 GTATGTGTACCTAAGGGAGTAGTGAGTTCACTGCTTGTTATGTATCATTTCAGGAGTTACTGG 2340

Qy 2341 ATTTGTTGTTCTTTCAGAAATACTAAAATTATTGTGTAGAAAGAAAACTGGAAA 2400
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 2341 ATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAAATTATTGTGTAGAAAGAAAACTGGAAA 2400

Qy 2401 CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCCGAATTCATGTGGTGCTTTTCACTT 2460
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 2401 CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCCGAATTCATGTGGTGCTTTTCACTT 2460

Qy 2461 GTGTTTTAAAATAAGGATTTTTCTCTTAATTCTCCCCCTTAGTTTGTGTGAGAAACATCTCAA 2520
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 2461 GTGTTTTAAAATAAGGATTTTTCTCTTAATTCTCCCCCTTAGTTTGTGTGAGAAACATCTCAA 2520

Qy 2521 TAAAGTGCTTTAAAATAAAATCTGAGCAATGCAGAGACACATCTCAA 2580
          ||||||||||||||||||||||||||||||||||||||||||||||
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Qy 2581 AAAAAAAAA 2589
          |||||||||
Db 2581 AAAAAAAAA 2589

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Qy   1081 ATTATGTGGGTCGCAATGATGATGATGTCAAATGCTTTTGTTGTGATGGTGGCTTGAGGTGTGTT 1140
Db   1081 GGGAAATCTGGGAGATGGATGAATCCAAGTGGTTTGTTTTCCAAGGTGTGTGAGTTCT 1200
Qy   1141 GGGGAAATCTGGGAGATGGATGAATCCAAGTGGTTTGTTTTCCAAGGTGTGTGAGTTCT 1200
Db   1141 GGGGAAATCTGGGAGATGGATGAATCCAAGTGGTTTGTTTTCCAAGGTGTGTGAGTTCT 1200
Qy   1201 TGATACGGAATGAAAAGGCCAAGAGAGTTGTTGGATTGAGAGTAGATATCCTCATCTTC 1260
Db   1201 TGATACGGAATGAAAAGGCCAAGAGAGTTGTTGGATTGAGAGTAGATATCCTCATCTTC 1260
Qy   1261 TTGAACAGCTGTTGTGTTCAACTTCAGATACCACTGGAGAAGCAAAATGCTGACCCACCAATTA 1320
Db   1261 TTGAACAGCTGTGTTCAACTTCAGATACCACTGGAGAAGCAAAATGCTGACCCACCAATTA 1320
Qy   1321 TTCATTTTGGGACCTGGAGAAAGTTCTTCAGAAGATGGCTGTCATGAATGAATACACCTGTGG 1380
Db   1321 TTCATTTTGGGACCTGGAGAAAGTTCTTCAGAAGATGGCTGTCATGAATGAATACACCTGTGG 1380
Qy   1381 TTAAAATCTGCCTTGGAAATGGGCTTTTAATACGAGACCTGGTGAAACAAACAGTTCAAAGTA 1440
Db   1381 TTAAAATCTGCCTTGGAAATGGGCTTTTAATACGAGACCTGGTGAAACAAACAGTTCAAAGTA 1440
Qy   1441 AAATCCTGACAACTGGAGAGAACTATATAAAACAGTTAATGATTATTTGGTGTCAGCACTTCTAA 1500
Db   1441 AAATCCTGACAACTGGAGAGAACTATATAAAACAGTTAATGATTATTTGGTGTCAGCACTTCTAA 1500
Qy   1501 ATGCTGAAGATGAAAAAAGAGAGGAAAACAAGCTGAAGAAATGGCATCAG 1560
Db   1501 ATGCTGAAGATGAAAAAAGAGAGGAAAACAAGCTGAAGAAATGGCATCAG 1560
Qy   1561 ATGATTTGTCATTAATTCGGAAGAACAGAAATGGCTCTCTTTCAACAATTGACATGTGTGC 1620
Db   1561 ATGATTTGTCATTAATTCGGAAGAACAGAAATGGCTCTCTTTCAACAATTGACATGTGTGC 1620
Qy   1621 TTCCTATCCTGGATAATCTTTTTAAAGGCCAATGTAATTAATAAACAGAACATGATAATTA 1680
Db   1621 TTCCTATCCTGGATAATCTTTTTAAAGGCCAATGTAATTAATAAACAGAACATGATAATTA 1680
Qy   1681 TTAAACAAAAAACACAGATACCTTTACAAGCGAGAGAACTGATTGAATGAATACCATTTGGTTA 1740
Db   1681 TTAAACAAAAAACACAGATACCTTTACAAGCGAGAGAACTGATTGAATGAATACCATTTGGTTA 1740
Qy   1741 AAGGAAAATGCTGGGCGCCAACATCTTCAAAAACTGTCTAAAAGAAAATTGACTCTACATTGT 1800
Db   1741 AAGGAAAATGCTGGGCGCCAACATCTTCAAAAACTGTCTAAAAGAAAATTGACTCTACATTGT 1800
Qy   1801 ATAAGAACTTATTTGTGGATAAGAATATGAAGTATATATTCCAACAGAAGATGTTTCAGGTC 1860
Db   1801 ATAAGAACTTATTTGTGGATAAGAATATGAAGTATATATTCCAACAGAAGATGTTTCAGGTC 1860
Qy   1861 TGTCACTCGGAAGAACAATTGAAGGAGGTTGCAAGAGAACGAACTTGTATAAAGTGTGTATGG 1920
Db   1861 TGTCACTCGGAAGAACAATTGAAGGAGGTTGCAAGAGAACGAACTTGTATAAAGTGTGTATGG 1920
Qy   1921 ACAAAGAGTTTCTGTGTATTATTCCTTGTGGTGGTCATCTGGTAGTAGTGCCAGGAAATGTG 1980
Db   1921 ACAAAGAGTTTCTGTGTATTATTCCTTGTGGTGGTCATCTGGTAGTAGTGCCAGGAAATGTG 1980
Qy   1981 CCCCTTCCTAAGAAAAATGCCCTATTTGCAGGGGTATAATCAAAGGGTACCTGTTCGGTACAT 2040
Db   1981 CCCCTTCCTAAGAAAAATGCCCTATTTGCAGGGGTATAATCAAAGGGTACCTGTTCGGTACAT 2040
Qy   2041 TTCTCTCTTAAAGAAAAATAGTCTATATATTTAAACCTGGCATAAAAAGGTCTTTAAAATATT 2100
Db   2041 TTCTCTCTTAAAGAAAAATAGTCTATATATTTAAACCTGGCATAAAAAGGTCTTTAAAATATT 2100
Qy   2101 GTTGAACAACTTGAAGGCCATCTAAAGTAAAAAAGGGAATTATGAGTTTTTTCAAATTAGTAACA 2160
Db   2101 GTTGAACAACTTGAAGGCCATCTAAAGTAAAAAAGGGAATTATGAGTTTTTTCAAATTAGTAACA 2160
Qy   2161 TTCATGTTCTAGTGTCTGCTTTGGTACTAATAATCTTGTTTCTGAAAAAGAATGGGTATCATATA 2220
Db   2161 TTCATGTTCTAGTGTCTGCTTTGGTACTAATAATCTTGTTTCTGAAAAAGAATGGGTATCATATA 2220
Qy   2221 TTTAATCTTAATCTGTTTTATTTACAAGGGAAGAATTAGTTGTTGGTGAACTAATATTAGTTAT 2280
Db   2221 TTTAATCTTAATCTGTTTTATTTACAAGGGAAGAATTAGTTGTTGGTGAACTAATATTAGTTAT 2280
Qy   2281 GTATGTGTTACCTTAAGGGAGGAGTAGTGTCACTGCTTGTTTATGCATCATTTCAGGAGTTACTGG 2340
Db   2281 GTATGTGTTACCTTAAGGGAGGAGTAGTGTCACTGCTTGTTTATGCATCATTTCAGGAGTTACTGG 2340
Qy   2341 ATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAATATAGTGTAGAAAAGAAACATGGAAA 2400
Db   2341 ATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAAATTAATAGTGTAGAAAAGAAACTGGAAA 2400
Qy   2401 CCAGGAACTCTGGAGTTCATCAGAGTTATGGGTGCCGAATTGTCTTTGGGTGCTTTCACTT 2460
Db   2401 CCAGGAACTCTGGAGTTCATCAGAGTTATGGGTGCCGAATTGTCTTTGGGTGCTTTCACTT 2460
Qy   2461 GTGTTTTAAAATAAGGATTTTCTCTTATTTCTCCCCCTAGTTTGTGAGAAACATCTCAA 2520
Db   2461 GTGTTTTAAAATAAGGATTTTCTCTTATTTCTCCCCCTAGTTTGTGAGAAACATCTCAA 2520
Qy   2521 TAAAGTGCTTTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2580
Db   2521 TAAAGTGCTTTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2580
Qy   2581 AAAAAAAAA 2589
Db   2581 AAAAAAAAA 2589
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RESULT 3

US-09-205-204-1

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;   Sequence 1, Application US/09205204
;   Patent No. 5958772
;   GENERAL INFORMATION:
;   APPLICANT: C. Frank Bennett
;   APPLICANT: Elizabeth J. Ackerman
;   APPLICANT: Lex M. Cowsert
;   TITLE OF INVENTION: ANTISENSE MODULATION OF CELLULAR INHIBITOR OF APOPTOSIS-1 EXPRES
;   FILE REFERENCE: RTS-0020
;   CURRENT APPLICATION NUMBER: US/09/205,204
;   CURRENT FILING DATE: 1998-12-03
;   NUMBER OF SEQ ID NOS: 47
;   SEQ ID NO 1
;   LENGTH: 3532
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (1160)..(3016)
US-09-205-204-1
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Query Match 97.9%; Score 2535.8; DB 2; Length 3532;
Best Local Similarity 99.7%;
Matches 2540; Conservative 0; Mismatches 7; Indels 0; Gaps 0;
Pred. No. 0;

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Qy      1 TCTAAGTAGTATCTTGAAATTCGAGAGATACTCATCCTACCTGAATATAAAACTGAGAT 60
Db    966 TCTAAGTAGTATCTTGAAATTCGAGAGATACTCATCCTACCTGAATATAAAACTGAGAT 1025
Qy     61 AAATCCAGTAAAGAAAAGTGTAGTAAAATTCTACATAAGAGTCTATAAGATTCTTTTTTGG 120
Db   1026 AAATCCAGTAAAGAAAAGTGTAGTAAAATTCTACATAAGAGTCTATAAGATTCTTTTTTGG 1085
Qy    121 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAATGTTTTAGCTATCAAAACAGC 180
Db   1086 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAATGTTTTAGCTATCAAAACAGT 1145
Qy    181 ACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAGAGGTCCCTCGTATC 240
Db   1146 ACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAGAGACTTTTCCCAGGTCCCTCGTATC 1205
Qy    241 AAAACATTAAGAGTATAATGGAAGATAGCACGGATCTTGTCAGATTGGACAAACAGCAACA 300
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                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1206 AAAACATTAAGAGTATAATGGAAGATAGCACGATCTTGTCAGATTGGACAAACAGCAACA 1265

Qy         301 AACAAAAAATGAAGTATGACTTTTCCTGTGAACTCTACAGAATGTCTACATATTCAACTT 360
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1266 AACAAAAAATGAAGTATGACTTTTCCTGTGAACTCTACAGAATGTCTACATATTCAACTT 1325

Qy         361 TCCCCGCCGGGGTGCCTGTCTCAGAAAGGAGTCTTGCTCGTGCTGGTTTTTATTATACTG 420
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1326 TCCCCGCCGGGGTGCCTGTCTCAGAAAGGAGTCTTGCTCGTGCTGGTTTTTATTATACTG 1385

Qy         421 GTGTGAATGACAAGGTCAAATGCTTCTGTTGTGGCCTGATGCTGGATAACTGGAAACTAG 480
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1386 GTGTGAATGACAAGGTCAAATGCTTCTGTTGTGGCCTGATGCTGGATAACTGGAAACTAG 1445

Qy         481 GAGACAGTCCTATTCAAAAGCATAAACAGCTATATCCTAGCTGTAGCTTTATTCAGAATC 540
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Db        1446 GAGACAGTCCTATTCAAAAGCATAAACAGCTATATCCTAGCTGTAGCTTTATTCAGAATC 1505

Qy         541 TGGTTTCAGCTAGTCTGGGATCCACCTCTAAGAATACGTCTCCAATGAGAAACAGTTTTG 600
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Db        1506 TGGTTTCAGCTAGTCTGGGATCCACCTCTAAGAATACGTCTCCAATGAGAAACAGTTTTG 1565

Qy         601 CACATTCATTATCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC 660
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Db        1566 CACATTCATTATCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC 1625

Qy         661 TTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGACTAACC 720
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1626 TTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGACTAACC 1685

Qy         721 CCTACAGTTATGCAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCATATGTGGCCAT 780
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Db        1686 CCTACAGTTATGCAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCATATGTGGCCAT 1745

Qy         781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATAGGACCTGGAG 840
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Db        1746 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATAGGACCTGGAG 1805

Qy         841 ATAGGGTAGCCTGCTTTGCCTGTGGTGGGAAGCTCAGTAACTGGGAACCAAAGGATGATG 900
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1806 ATAGGGTAGCCTGCTTTGCCTGTGGTGGGAAGCTCAGTAACTGGGAACCAAAGGATGATG 1865

Qy         901 CTATGTCAGAACACCGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAATTCTCTAGAAA 960
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Db        1866 CTATGTCAGAACACCGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAATTCTCTAGAAA 1925

Qy         961 CTCTGAGGTTTAGCATTTCAAATCTGAGCATGCAGACACATGCAGCTCGAATGAGAACAT 1020
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Db        1926 CTCTGAGGTTTAGCATTTCAAATCTGAGCATGCAGACACATGCAGCTCGAATGAGAACAT 1985

Qy        1021 TTATGTACTGGCCATCTAGTGTTCCAGTTCAGCCTGAGCAGCTTGCAAGTGCTGGTTTTT 1080
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Db        1986 TTATGTACTGGCCATCTAGTGTTCCAGTTCAGCCTGAGCAGCTTGCAAGTGCTGGTTTTT 2045

Qy        1081 ATTATGTGGGTCGCAATGATGATGTCAAATGCTTTTGTTGTGATGGTGGCTTGAGGTGTT 1140
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Db        2046 ATTATGTGGGTCGCAATGATGATGTCAAATGCTTTTGTTGTGATGGTGGCTTGAGGTGTT 2105

Qy        1141 GGGAATCTGGAGATGATCCATGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 1200
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Db        2106 GGGAATCTGGAGATGATCCATGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 2165

Qy        1201 TGATACGAATGAAAGGCCAAGAGTTTGTTGATGAGATTCAAGGTAGATATCCTCATCTTC 1260
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Db        2166 TGATACGAATGAAAGGCCAAGAGTTTGTTGATGAGATTCAAGGTAGATATCCTCATCTTC 2225

Qy        1261 TTGAACAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCCACCAATTA 1320
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        2226 TTGAACAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCCACCAATTA 2285

Qy        1321 TTCATTTTGGACCTGGAGAAAGTTCTTCAGAAGATGCTGTCATGATGAATACACCTGTGG 1380
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        2286 TTCATTTTGGACCTGGAGAAAGTTCTTCAGAAGATGCTGTCATGATGAATACACCTGTGG 2345

Qy        1381 TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGACCTGGTGAAACAAACAGTTCAAAGTA 1440
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Db        2346 TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGACCTGGTGAAACAAACAGTTCAAAGTA 2405

Qy        1441 AAATCCTGACAACTGGAGAGAACTATAAAACAGTTAATGATATTGTGTCAGCACTTCTAA 1500
                |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        2406 AAATCCTGACAACTGGAGAGAACTATAAAACAGTTAATGATATTGTGTCAGCACTTCTTA 2465

Qy        1501 ATGCTGAAGATGAAAAAAGAGAGGAGGAGAAGGAAAAACAAGCTGAAGAAATGGCATCAG 1560
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Db        2466 ATGCTGAAGATGAAAAAAGAGAAGAGGAGAAGGAAAAACAAGCTGAAGAAATGGCATCAG 2525

Qy        1561 ATGATTTGTCATTAATTCGGAAGAACAGAATGGCTCTCTTTCAACAATTGACATGTGTGC 1620
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Db        2526 ATGATTTGTCATTAATTCGGAAGAACAGAATGGCTCTCTTTCAACAATTGACATGTGTGC 2585

Qy        1621 TTCCTATCCTGGATAATCTTTTAAAGGCCAATGTAATTAATAAACAGGAACATGATATTA 1680
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Db        2586 TTCCTATCCTGGATAATCTTTTAAAGGCCAATGTAATTAATAAACAGGAACATGATATTA 2645

Qy        1681 TTAAACAAAAAACACAGATACCTTTACAAGCGAGAGAACTGATTGATACCATTTTGGTTA 1740
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Db        2646 TTAAACAAAAAACACAGATACCTTTACAAGCGAGAGAACTGATTGATACCATTTTGGTTA 2705

Qy        1741 AAGGAAATGCTGCGGCCAACATCTTCAAAAACTGTCTAAAAGAAATTGACTCTACATTGT 1800
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Db        2706 AAGGAAATGCTGCGGCCAACATCTTCAAAAACTGTCTAAAAGAAATTGACTCTACATTGT 2765

Qy        1801 ATAAGAACTTATTTGTGGATAAGAATATGAAGTATATTCCAACAGAAGATGTTTCAGGTC 1860
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Db        2766 ATAAGAACTTATTTGTGGATAAGAATATGAAGTATATTCCAACAGAAGATGTTTCAGGTC 2825

Qy        1861 TGTCACTGGAAGAACAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTGTATGG 1920
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Db        2826 TGTCACTGGAAGAACAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTGTATGG 2885

Qy        1921 ACAAAGAAGTTTCTGTTGTATTTATTCCTTGTGGTCATCTGGTAGTATGCCAGGAATGTG 1980
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        2886 ACAAAGAAGTTTCTGTTGTATTTATTCCTTGTGGTCATCTGGTAGTATGCCAGGAATGTG 2945

Qy        1981 CCCCTTCTCTAAGAAAATGCCCTATTTGCAGGGGTATAATCAAGGGTACTGTTCGTACAT 2040
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Db        2946 CCCCTTCTCTAAGAAAATGCCCTATTTGCAGGGGTATAATCAAGGGTACTGTTCGTACAT 3005

Qy        2041 TTCTCTCTTAAAGAAAAATAGTCTATATTTTAACCTGCATAAAAAGGTCTTTAAAATATT 2100
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Db        3006 TTCTCTCTTAAAGAAAAATAGTCTATATTTTAACCTGCATAAAAAGGTCTTTAAAATATT 3065

Qy        2101 GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGGAATTATGAGTTTTTCAATTAGTAACA 2160
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Db        3066 GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGGAATTATGAGTTTTTCAATTAGTAACA 3125

Qy        2161 TTCATGTTCTAGTCTGCTTTGGTACTAATAATCTTGTTTCTGAAAAGATGGTATCATATA 2220
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Db        3126 TTCATGTTCTAGTCTGCTTTGGTACTAATAATCTTGTTTCTGAAAAGATGGTATCATATA 3185

Qy        2221 TTTAATCTTAATCTGTTTATTTACAAGGGAAGATTTATGTTTGGTGAACTATATTAGTAT 2280
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Db        3186 TTTAATCTTAATCTGTTTATTTACAAGGGAAGATTTATGTTTGGTGAACTATATTAGTAT 3245

Qy        2281 GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTATGCATCATTTCAGGAGTTACTGG 2340
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Db        3246 GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTATGCATCATTTCAGGAGTTACTGG 3305

Qy        2341 ATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAATTATAGTGTAGAAAAGAACTGGAAA 2400
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Db        3306 ATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAATTATAGTGTAGAAAAGAACTGGAAA 3365

Qy        2401 CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCCGAATTGTCTTTGGTGCTTTTCACTT 2460
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Db        3366 CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCCGAATTGTCTTTGGTGCTTTTCACTT 3425
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RESULT 4

US-09-212-971-7

; Sequence 7, Application US/09212971B
; Patent No. 6107041
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; FILE REFERENCE: 07891/009002
; CURRENT APPLICATION NUMBER: US/09/212,971B
; CURRENT FILING DATE: 1998-12-16
; EARLIER APPLICATION NUMBER: 60/017,354
; EARLIER FILING DATE: 1996-04-26
; EARLIER APPLICATION NUMBER: 60/030,590
; EARLIER FILING DATE: 1996-11-14
; EARLIER APPLICATION NUMBER: 08/800,929
; EARLIER FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 7
; LENGTH: 3732
; TYPE: DNA
; ORGANISM: Homo sapiens

US-09-212-971-7

Query Match   97.9%;   Score 2535.8;   DB 3;   Length 3732;
Best Local Similarity   99.7%;   Pred. No.: 0;
Matches 2540;   Conservative   0;   Mismatches   7;   Indels   0;   Gaps   0;

Qy   1 TCTAAGTAGTATCTTGGAAATTCAGAGAGAATACTCATCCTACCTGAATATAAAACTGAGAT 60
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Db   1245 TCTAAGTAGTATCTTGGTAATTCAGAGAGATACTCATCCTACCTGAATATAAAACTGAGAT 1186

Qy   61 AAATCCAGTAAAGAAAGTAGTAGTAAATTCTACATAAGAGTCTATCATACATTGATTGATT 120
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1305 AAATCCAGTAAAGAAAGTAGTAGTAAATTCTACATAAGAGTCTATCATACATTGATTGATT 1246

Qy   121 TGGTAAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAATGTTTAGCTATCAAACAGC 180
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Db   1365 TGGTAAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAATGTTTAGCTATCAAACAGT 1306

Qy   181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCCAAAGAGACTTTTCCCAGGTCCCTCGTATC 240
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Db   1425 ACTGTCACCTACTCATGCACAAAACTGCCTCCCCAAAGACTTTTCCCAGGTCCCTCGTATC 1366

Qy   241 AAAACATTAAGAGTATAATAGGAAGATAGCACGAATCTTGTCAGATTGGCAAACAGCAACA 300
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Db   1485 AAAACATTAAGAGTATAATGGAAGATAGCACGAATCTTGTCAGATTGGCAAACAGCAACA 1426

Qy   301 AACAAAAAAATGAAGTAGTATGACTTTTTCCTGTGAAGAATGTCTACATAATTCAACTT 360
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Db   1545 AACAAAAAAATGAAGTATGACTTTTTCCTGTGAAGAATGTCTACATAATTCAACTT 1486

Qy   361 TCCCCGCGCCGGGGGTGCCTGCTTGTTGCTCGGTTCTGGTCGTGGTTTTTTATTATATACTG 420
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Db   1605 TCCCGCGCCGGGGGTGCCTGCTTGTTGCTCGGTTCTGGTCGTGGTTTTTTATTATATACTG 1546

Qy   421 GTGTGAATGACAAAGGTCAAAATGCTTCTGTTGGTTGTGGGCCTGGATAAACTGAAACTAG 480
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Db   1665 GTGTGAATGACAAAGGTCAAAATGCTTCTGTTGGTTGTGGGCCTGGATAAACTGAAACTAG 1606

Qy   481 GAGACAGTCCTATTCAAAGACATAAAACAGCTATATCCTAGCTGTTATTCAGAATC 540
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Db   1725 GAGACAGTCCTATTCAAAGACATAAAACAGCTATATCCTAGCTGTTATTCAGAATC 1666

Qy   541 TGGTTTCAGCTAGTCTGGGATCCACCCTCTAAGAATATAGCTCTCCAATGAGAAACAGTTTTG 600
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Db   1785 TGGTTTCAGCTAGTCTGGGATCCACCCTCTAAGAATATAGCTCTCCAATGAGAAACAGTTTTG 1726

Qy   601 CACATTCATTATCTCCCACCTTGGAACATGAGACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC 660
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Db   1845 CACATTCATTATCTCCCACCTTGGAACATGAGACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC 1786

Qy   661 TTTCTCCAAACCCTCTTAATTCTAGGAGCAGTTGAAGACATCTTCATCGAGGACTAACC 720
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Db   1905 TTTCTCCAAACCCTCTTAATTCTAGGAGCAGTTGAAGACATCTTCATCGAGGACTAACC 1846

Qy   721 CCTACAGTTATGCAAATGACTACTGAAGAAGGCCAGATTTTCTTACCTACCATAGTTGGCCAT 780
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Db   1965 CCTACAGTTATGCAAATGACTACTGAAGAAGGCCAGATTTTCTTACCTACCATAGTTGGCCAT 1906

Qy   781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTATTATATATAGGACCTGGAG 840
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Db   2025 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTATTATATAGGACCTGGAG 1966

Qy   841 ATAGGGTAGCCTGCTTGTGCCTGTTGGTGGGGAAGCTCACTGTAACTCGGGACACAAAGAGGATGATG 900
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Db   2085 ATAGGGTAGCCTGCTTGTGCCTGTTGGTGGGGAAGCTCAGTGAACCAAAGAGGATGATG 2026

Qy   901 CTATGTCAGAACACCGGAGGGCCATTTCCCAACTGTCCAACTGTGTCCATTTTTTGGAAAAATTCTTCTAGAA 960
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Db   2145 CTAGTCAGAACACCGGAGGGCCATTTTCCCAACTGTCCATTTTTTGGAAAATTCTCTAGAAA 2086

Qy   961 CTCTGAGGTTTAGCATTTCAAATCTCAGCATCTCAGAGACAACATGCAGCTCGAATGAGAACAT 1020
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Db   2205 CTCTGAGGTTTAGCATTTCAAATCTCAGCATCTCAGAGACAACATGCAGCTCGAATGAGAACAT 2146

Qy   1021 TTATGTACTGGCCATCTAGTGTGTTCCAGTTCAGCCTGAGCGCTGCAAGTGCTGGTTTTTT 1080
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Db   2265 TTAAGTACTGGCCATCTAGTGTGTTCCAGTTCAGCCTGAGCGCTGCAAGTGCTGGTTTTT 2206

Qy   1081 ATTATGTGGGTCGCAATGATCAATGATGATTCTTTGGTTGTTTGGATGGTGGTTGAGGTGTT 1140
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Db   2325 ATTATGTGGGTCGCAATGATCAATGATGATTCTTTTGTTGTTTGGATGGTGGTTGAGGTGTT 1140

Qy   1141 GGGGAATCTGGAGATGAATCCATGGGTAGGAACATGCCAAGTGGTGAGTTCT 1200
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Db   2385 GGGGAATCTGGAGATGAATCCATGGGTAGGAACATGCCAAGTGGTGAGTTCT 2326

Qy   1201 TGATACGAAATGAAAAGGCCAAGAGTTTGTTGATGAGATTCAAAGGTAGAGATATCCTCAACCTTC 1260
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Db   2445 TGATACGAAATGAAAAGGCCAAGAGTTTGTTGATGAGATTCAAAGGTAGAGATATCCTCAACCTTC 1260

Qy   1261 TTGAACAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAGAAAATGGTGACCCACCAATTA 1320
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Db   2505 TTGAACAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCCACCAATTA 1320

Qy   1321 TTCATTTTGGACCTGGAGAAATGTTCTTCGAGAGATGCTTGTCATGATGAATACACCTGTGG 1380
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Db   2565 TTCATTTTGGACCTGGAGAAATGTTCTTCGAGAGATGCTTGTCATGATGAATACACCTGTGG 1380

Qy   1381 TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGAGACCTGGGTTGAAAACAGTTCAAAGTA 1440
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2625 TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGAGACCTGGGTTGAAAACAGTTCAAAGTA 1440

Qy   1441 AAATCCTGACAACTGGAGAGAACTATAAAACAGTTAATGATATTGTGTCAGCAC 1500
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2685 AAATCCTGACAACTGGAGAGAACTATAAAAACAGTTAATGATATTGTGTCAGCACTTCTTA 2626

Qy   1501 ATGCTGAAGATGAAAAAAGAGAGGAGAAAACAAAGCTGAAGAAATGGCATCAG 1560
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1560

RESULT 5

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Qy   3706 TAAAGTGCTTTAAAAAGAAAAAAAAAAAAA 3732
          |||||||||||||||||
Db   2521 TAAAGTGCTTTAAAAAGAAAAAAAAAAAA 2547
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US-08-800-929A-7
; Sequence 7, Application US/08800929A
; Patent No. 6133437
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF
; TITLE OF INVENTION: IAPS AND NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESS:
;   ADDRESSEE: Clark & Elbing LLP
;   STREET: 176 Federal Street
;   CITY: Boston
;   STATE: MA
;   COUNTRY: USA
;   ZIP: 02110
; COMPUTER READABLE FORM:
;   MEDIUM TYPE: Diskette
;   COMPUTER: IBM Compatible
;   OPERATING SYSTEM: DOS
;   SOFTWARE: FastSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08/800,929A
;   FILING DATE: 13-FEB-1997
;   CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: 60/030,590
;   FILING DATE: 14-NOV-1996
;   APPLICATION NUMBER: 60/017,354
;   FILING DATE: 26-APR-1996
; ATTORNEY/AGENT INFORMATION:
;   NAME: Bieker-Brady, Kristina
;   REGISTRATION NUMBER:
;   REFERENCE/DOCKET NUMBER: 07891/009001
; TELECOMMUNICATION INFORMATION:
;   TELEPHONE: 617-428-0200
;   TELEFAX: 617-428-7045
;   TELEX:
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
;   LENGTH: 3732 base pairs
;   TYPE: nucleic acid
;   STRANDEDNESS: single
;   TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-800-929A-7

Query Match   97.9%; Score 2535.8; DB 3; Length 3732;
Best Local Similarity   99.7%; Pred. No. 0; Mismatches 7; Indels 0; Gaps 0;
Matches 2540; Conservative 0; Mismatches 7; Indels 0; Gaps 0;
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Qy      1 TCTAAGTAGTATCTTGGAAATTCAGAGAGATAGTACTCATCCTACCTGAAATATAAACTGAGAT 60
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Db   1186 TCTAAGTAGTATCTTGGTATTCAGAGAGATAGTACTCATCCTACCTGAAATATAAACTGAGAT 1245

Qy     61 AAATCCAGTAAAGAAAGTGTAGTAGTTAAATTCTACATTAAGAGTCTATCATTGATTTCTTTTGG 120
          ||||||||||||||||||||||||||||||||||||||||||
Db   1246 AAATCCAGTAAAGAAAGTGTAGTAGTTAAATTCTACATTAAGAGTCTATCATTGATTTCTTTTG 1305

Qy    121 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAAATGTTTTAGTTATCAAACAGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1306 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAAATGTTTTAGTTATCAAACAGT 1365

Qy    181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTCGTATC 240
          |||||||||||||||||||||||||||||||||||||||||||||||
Db   1366 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCATGTGAAATGTTTTAGCTTATCAAACAGT 1425

Qy    241 AAAACATTAAGAGTATAATGAAGATAGCACGATCTTGTCAGATTGGACAAACAGCAACA 300
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Qy   1561 ATGATTTGTCATTAATTCGGAGAAGACAGAATGGCTCTCTTCTTCAACAATTGACATGTGTGC 1620
          |||||||||||||||||||||||||||||||||
Db   2686 ATGATTTGTCATTAATTCGGAGAAGACAGAATGGCTCTCTTCTTCAACAATTGACATGTGTGC 2745
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... (alignment continues)

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Db  1426 AAAAACATTAAGAGTATAATGTGGAAGATAGCACGATCTTTGTCAGATTGGACAACAGCAACA          1485
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Qy   301 AACAAAAAATGAAGTAATGACTTTTCCTGTGCTGTGAAACTTCACAGAAATGCTACATATTCAACTT          360
Db  1486 AACAAAAAATGAAGTAATGACTTTTCCTGTGCTGTGAAACTTCACAGAAATGCTACATATTCAACTT          1545
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy   361 TCCCCCCGCGGGGGGTGCCTGTCTCAGAAAAGAGAGTCTTTGCCTCTGGTGGTTTTTTATTATACTG          420
Db  1546 TCCCCCCGCGGGGGGTGCCTGTCTCAGAAAAGAGAGTCTTTGCCTCTGGTGGTTTTTTATTATACTG          1605
Qy   421 GTGTGAAATGACAAGGTCAAAATGCTTTTCTTGTTGTTGTGGGGCCTTGGATAACTGGAAACTAG          480
Db  1606 GTGTGAAATGACAAGGTCAAAATGCTTTTCTTGTTGTTGTGGGGCCTTGGATAACTGGAAACTAG          1665
Qy   481 GAGACAGTCCTATTCTCAAAAAGCATAAAACAGTATATCCTAGCTGTAGCTTTACATATTCAGAAATC          540
Db  1666 GAGACAGTCCTATTCTCAAAAAGCATAAAACAGTATATCCTAGCTGTAGCTTTACATATTCAGAAATC          1725
Qy   541 TGGGTTTCAGCTAGTCTGGGATCCACCCTTAAGAATAGTTCTCCAATGAGAAACAGTTTTG          600
Db  1726 TGGGTTTCAGCTAGTCTGGGATCCACCCTTAAGAATAGTTCTCCAATGAGAAACAGTTTTG          1785
Qy   601 CACATTCATTTACTCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGAGTTCTTTACTCCAGCC          660
Db  1786 CACATTCATTTACTCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGAGTTCTTTACTCCAGCC          1845
Qy   661 TTTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGACTAACC          720
Db  1846 TTTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGACTAACC          1905
Qy   721 CCTACAGTTATGCAATGAGTACTGAAGAAGCCAGATTTTCTTACCTACCATATGTTGACCAT          780
Db  1906 CCTACAGTTATGCAATGAGTACTGAAGAAGCCAGATTTTCTTACCTACCATATGTTGACCAT          1965
Qy   781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTTATTATAGGACCTGGAG          840
Db  1966 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTTATTATAGGACCTGGAG          2025
Qy   841 ATAGGGGTAGCCCTCCTTTGCCTGTGGTGGGAAGCTCAGTAACTGGGAACCAAAAGGATGATG          900
Db  2026 ATAGGGGTAGCCCTCCTTTGCCTGTGGTGGGAAGCTCAGTAACTGGGAACCAAAAGGATGATG          2085
Qy   901 CTATGTCAGAACACCGGAGGCCAATTTTCCCAAACTGTCCAATTTTTTGGAAAATTCTCTAGAA          960
Db  2086 CTATGTCAGAACACCGGAGGCCAATTTTCCCAAACTGTCCAATTTTTTGGAAAATTCTCTAGAA          2145
Qy  1021 TTATGTACTGGCCATCTACTGTGTTCCAGTTCAGCCTGAAGTGCTCTGGTTTTT          1080
Db  2206 TTATGTACTGGCCATCTACTGTGTTCCAGTTCAGCCTGAAGTGCTCTGGTTTTT          2265
Qy  2146 CTCTGAGGTTTAGCATTTCAAATCTGAGCAATGCAGACACAGTGAAATGAGAAACAT          2205
Db   961 CTCTGAGGTTTAGCATTTCAAATCTGAGCAATGCAGACACAGTGAAATGAGAAACAT          1020
Qy  2086 CTATGTCAGAACACCGGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAATTCTCTAGAAA          2145
Qy  1906 CCTACAGTTATGCAATGAGTACTGAAGAAGCCAGATTTTCTTACCTACCATATGTTGACCAT          1965
Db  1966 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTTATTATAGGACCTGGAG          2025
Qy  1846 TTTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGACTAACC          1905
Db  1801 ATAAGAAACTTATTTGTGGGATAAGAATATGAGGAGGTTGCAAGGATCAAAGATTGACTTACATTGT          1860
Qy  2026 ATAGGGGTAGCCTGCTTTGCCTGTGGGAAGCTCAGTAACTGGGAACCAAAAGGATGATG          2085
Qy  1861 TGTCACTGGAAGAACAATTGAGGAGGTTGCAAGAGAACGAACTTGTGAAGTGTATGGG          1920
Db  3046 TGTCACTGGAAGAACAATTGAGGAGGTTGCAAGAGAACGAACTTGTGAAGTGTATGGG          3105
Qy  1801 ATAAGAAACTTATTTGTGGGATAAGAATATGAGGAGGTTCATCGGTTCATCGGGAAATGTG          1860
Db  1921 ATAAGAAACTTATTTGTGGGATAAGAATATGAGGAGGTTCATCGGTTCATCGGGAAATGTG          1980
Qy   601 CACATTCATTTACTCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGAGTTCTTTACTCCAGCC          660
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Qy   2461 GTGTTTTAAAATAAGAGATTATTTTCTCTCCCCTAGTTTGTGAGAGAAACATCTCAA 2520
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3646 GTGTTTTAAAATAAGAGATTATTTTCTCTCCCCTAGTTTGTGAGAGAAACATCTCAA 3705

Qy   2521 TAAAGTGCTTTAAAAAAAAAAAAAA 2547
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Db   3706 TAAAGTGCTTTAAAAAGAAAAAAAAAAAAA 3732
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RESULT 6
US-09-617-053A-7
Sequence 7, Application US/09617053A
Patent No. 6300492
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009003
; CURRENT APPLICATION NUMBER: US/09/617,053A
; CURRENT FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 08/800,929
; PRIOR FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 7
; LENGTH: 3732
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-617-053A-7

Query Match   97.9%;   Score 2535.8;   DB 4;   Length 3732;
Best Local Similarity   99.7%;   Pred. No. 0;
Matches 2540;   Conservative   0;   Mismatches 7;   Indels 0;   Gaps 0;

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Qy      1 TCTAAGTAGTATCTTGGAAATTCAGAGAGAGATACTCATCCTCACCTGAATATAAACTGAGAT 60
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Db   1186 TCTAAGTAGTATCTTGGTAATTCAGAGAGAGATACTCATCCTCACCTGAATATAAACTGAGAT 1245

Qy     61 AAAATCCAGTAAAGAAAAGTGTAGTAGTAAATTCTACATAAAGACTGTCTATCTTTGATTTCTTTTTTTGG 120
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1246 AAAATCCAGTAAAGAAAAGTGTAGTAGTAAATTCTACATAAAGAGTCTATCATTGATTTCTTTTTTGG 1305

Qy    121 TGGTAAAAAATCTTAGTTCATGTGAAGAAAAATTCAGTGTGAAAGTTTTAAGCTATCAAACAGC 180
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1306 TGGTAAAAAATCTTAGTTCATGTGAAGAAAATTCATGTGAAAGTTTTAGCTATCAAACAGT 1365

Qy    181 ACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAGAGACTTTTCCCCAGGTTCCCTCGTATC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1366 ACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAGACTTTTCCCCAGGTTCCCTCGTATC 1425

Qy    241 AAAACATTAAGAGTATAAATGGAAGAAAGAACACAGARCTTGTCAGAATTGGACAACAGCAACA 300
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Db   1426 AAAACATTAAGAGTATAAATGGAAGAGAGAACACGATCTTGTGTCAGAATTGGACAAACAGCAACA 1485

Qy    301 AACAAAAAATGAAGTAAGTATGACTTTTCCGTGACTTACATCACATATTCAACTT 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1486 AACAAAAAATGAAGTAAGACTTTTCCGTGACTTACATCACATATTCAACTT 1545

Qy    361 TCCCCGCGCCGGGGGGTGCCTGTTCTCAGAAAGGAGTCTTGCTCGNGCTGGGTTTTTTTATTATACTG 420
          |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1546 TCCCGCGCCCGGGGTGCCTGTTCTCAGAATGGGCTGCTGGTTTTTTTATTATACTG 1605

Qy    421 GTGTGAAATGACAAGGTCAAAATGCTTCTCTGTTGTGTGGCCCTGATNTCTGGAACTAG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||
Db   1606 GTGTGAAATGACAAGGTCAAAATGCTTCTCGTTGTGTGGCCCTGATNTCTGGAAACTAG 1665

Qy    481 GAGACAGTCCTATTCAAAAAGCAATAAACAGCTATATATCCTAGCTGTAGCTTTATTCAGAAATC 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1666 GAGACAGTCCTATTCAAAAAGCAATAAACAGCTATATATCCTAGCTGTAGCTTTATTCAGAAATC 1725

Qy    541 TGGTTTCAGCTAGTCTGGGGATCCAACCTTCTAAAAGCTCCAAATGAGAAACAGTTTTG 600
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Qy    601 CACATTCAATTATCTCCCACCTTGGAACATTAGTAGCTTGTTTCAGTGTGGTTCTTACTCCAGCC 660
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Db   1786 CACATTCAATTATCTCCCACCTTGGAACATAGTAGCTTGTTTCAGTGTGGTTCTTACTCCAGCC 1845

Qy    661 TTTCTCCAAACCCTCTTAATTCTAGAGCAGTGAAGACATCTCTTCATCGAGGACTAAACC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1846 TTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGACTAAACC 1905

Qy    721 CCTACAGTTATGCAATGACGTACTGAAGAAGCCAGATTTCTTACCTACCATTATGTGGCCAT 780
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1906 CCTACAGTTATGCAATGACGTACTGAAGAAGCCAGATTTCTTACCTACCATTATGTGGCCAT 1965

Qy    781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTTATTATTATAGGACCTGGAG 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1966 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTTATTATATAGGACCTGGAG 2025

Qy    841 ATAGGGTAGCCCTGCTTTGCCTGTGCTTCAGTAGTGCGAAGCTCAGTAACTGGGAACCAAGATGATG 900
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2026 ATAGGGTAGCCCTGCTTTGCCTGTGTGGAAGCTCAGTAACTGGGAACCAAAGGATGATG 2085

Qy    901 CTATGTCAGAACACCCGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAAAAATTCTCTAGAAA 960
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Db   2086 CTATGTCAGAAACACCCGGAGGCCATTTTCCCCAACTGTCCATTTTTGGAAAAAAATTCTCTAGAAA 2145

Qy    961 CTCTGAGGTTTAGCATTTCAAAATCTGAGCATGCAGCACACATGCAGCTCGAAATGAGAACAT 1020
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2146 CTCTGAGGTTTAGCATTTCAAAATCTGAGCATGCAGCACATGCAGCTCGAAATGAGAACAT 2205

Qy   1021 TTATGTACTGGCCATCTAGTGTTCCAGTTCAGCCTGAGCAAGTGCTTGGTTTTT 1080
          |||||||||||||||||||||||||||||||||||||||||||||||
Db   2206 TTATGTACTGGCCCATCTAGTGTTCCAGGTTCAGCCTGAGCAAGTGCTTGGTTTTT 2265

Qy   1081 ATTATGTNGGGGTCCGCAAATGATGAATGTCAAATGCTTTGTTGTGATGGTGGCTTGAGGTGTT 1140
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2266 ATTATGTGGGGTCGCAAATGATGAATGTCAAATCCTTTGTTGTGATGGTGGCTTGAGGTGTT 2325

Qy   1141 GGGAATCTGGAGATGATCCATGGGTAGAACATGCCAAGGTGGTGAGTTCT 1200
          |||||||||||||||||||||||||||||||||||||||||||||||||
Db   2326 GGGAATCTGGAGATGATCCATGGGTAGAACATGCCAAGGTGTGAGTTCT 2385

Qy   1201 TGATACGGAATGAAAGGCCAAGAGTTGTTGTGATGAGTTCAGGTAGATATCCTCATCTTC 1260
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Db   2386 TGATACGGAATGAAAGGCCAAGAGTTTGTTGAGATTTCAAGGTAGATATCCTCATCTTC 2445

Qy   1261 TTGAAACAGCTGTGTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCCACCAACAATTA 1320
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Db   2446 TTGAAACAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAGTGCTGACCCACCAAATTA 2505

Qy   1321 TTCATTTTGGACCTGGAGAAAGTTCTTCAGAAGAGATGGCTGTGTCATGATAGAATACACCTGTGG 1380
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Db   2506 TTCATTTTGGACCTGGAGAAAGTTCTTCAGAAGAGATGCTGTGTCATGATAGAATACACCTGTGG 2565

Qy   1381 TTAAATCTGCCTTGGAAATGGGCTTTAATACGAAGACCTGGTGAAAACAAACAGTTCAAAGTA 1440
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Db   2566 TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGAGACCTGGTGAAAACAAACAGTTCAAAGTA 2625

Qy   1441 AAATCCTGACAACTGGGAGAGAACTATTAAAACAGTTAATGATATTGTGTCAGCACTTCTTA 1500
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Db   2626 AAATCCTGACAACTGGGAGAGAACTATTAAAACAGTTAATGATATTGTGTCAGCACTTCTTA 2685

Qy   1501 ATGCTGAAGGATGAAAAAAGAGGAGGAAAAAAACAAGCTGAAGAAAATGGCATCAG 1560
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Db   2686 ATGCTGAAGGATGAAAAAAGAGGAGGAAAAAAACAAGCTGAAGAAAATGGCATCAG 2745

Qy   1561 ATGATTTGTCATTAATTCGGAAGAACAGAGAATGGCTCTCTTCAAACAATTGACATGTGTGC 1620
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Db   2686 ... 
```

APPLICANT: Korneluk, Robert G.
APPLICANT: Mackenzie, Alexander E.
APPLICANT: Baird, Stephen
TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS, PROBES, AND DETECTION METHODS
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/511,485
FILING DATE: 04-AUG-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 07540/002001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 2580 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
MOLECULE TYPE: DNA (genomic)
US-08-511-485-7

Query Match 97.1%; Score 2514.8; DB 2; Length 2580;
Best Local Similarity 99.4%; Pred. No. 0; Mismatches 14; Indels 0; Gaps 0;
Matches 2522; Conservative 0;

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Qy      1  TCTAAGTAGTATCTTGGAAATTCAGAGAGAGATACTCATCCTACCTGAATATAAACTGAGAT   60
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Db      1  TCTAAGTAGTATCTTGGTAAATTCAGAGAGATACTCATCCTACCTGAATATAAACTGAGAT   60

Qy     61  AAAATCCAGTAAAGAGAAAGTGTAGTAAAAATTCTACATAAGAGTCTATCATTGATTCTTTTTGG  120
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Db     44  TCTAAGTAGTATCTTGGTAAATTCAGAGAGATACTCATCCTACCTGAATATAAACTGAGAT  103

Qy    121  TGGGTAAAATCTTAGTTCATGTGAAGAAAATTCATGTGAAAGTTTTAGCTATCAAAACAGC  180
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    104  AAAATCCAGTAAAGAGAAAGTGTAGTAAAAATTCTACATAAGAGTCATTGATTTCTTTTTG  163

Qy    181  ACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAAGACTTTTCCCAGGTCCCTCGTATC  240
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Db    164  TGGGTGGAAATCTTAGTTCATGTGAAGAAAATTCATGTGAAAGTTTTAGCTATCAAAACAGT  223

Qy    241  AAAACATTAAGACTATAATATTGGACACAGACGATCTTGTCAGATTGGACAAAACAGCAACA  300
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    224  ACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAAGACTTTTCCCAGGTCCCTCGTATC  283

Qy    301  AACAAAAAATGAAGTATGAACTTTTCCCGTGGAAAGAGAGAGTCTCACACAGAATGTCTAACTT  360
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    284  AAAACATTAAGACTATAATTGGACACAGACGATCTTGTCAGATTGGACAAAACAGCAACA  343

Qy    361  TCCCCGCGCGGGGTGCCCTGTCTCAGAAAAGGAGTCTTCGTCGTGGTTTTTATTATACTG  420
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    344  AACAAAAAATGAAGTATGAACTTTTCCCGTGTGAAACTCTTACACAGAATGTCTAACTT  403

Qy    421  GTGTGAATGACAAGGTCAAAATGCTTCTGTTGTTGTGGCCTGGGATAACTGGAAACTGG  480
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    404  TCCCCGCGCGGGGTGCCCTGTCTCAGAAAAGGAGTCTTCGTCGTGGTTTTTATTATACTG  463

Db    421  GTGTGAATGACAAGGTCAAAATGCTTCTGTTGTTGTGGCCTGGGATAACTGGAAACTGG  480
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Qy   1621  TTCCTATCCTGGATAAATCTTTTAAAGGCCAATGTAATTAATAAACACGGAACATGATATTA  1680
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2746  ATGATTTGTTCATTAAATTCGGAAGAACAGACAGAATGGCTCTCTTTCAACAATTGACATGTGTC  2805

Qy   1681  TTAAACAAAAAAACACAGATACCTTTTACAACGCGAGAAC TGATTGATACCATTTTGGTTA  1740
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2806  TTCCTATCCTGGATAAATCTTTTAAAGGCCAATGTAATTAATAAACACGGAACATGATATTA  2865

Qy   1741  AAGGAAATGCTGCGGCCAACATCTTCAAAAAACTGTCTAAAAAGAAAATTGACTCTACATTGT  1800
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2866  TTAAACAAAAAAACACAGATACCTTTTACAACGCGAGAACTGATTGATACCATTTTGGTTA  2925

Qy   1801  ATAAGAACTTATTTGTGGATAAGAATATGAAGTATATTCCAACAGAGAAGATGTTTCAGGTC  1860
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2926  AAGGAAATGCTGCGGCCAACATCTTCAAAAAACTGTCTAAAAAGAAAATTGACTCTACATTGT  2985

Qy   1861  TGTCACTGGAAGACAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTGTATGG  1920
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2986  ATAAGAACTTATTTGTGGATAAGAATATGAAGTATATTCCAACAGAGAAGATGTTTCAGGTC  3045

Qy   1921  ACAAAGAAGTTTCTGTTGTATTTATTTCCTTGTGGTCATCTGGTAGTAGTATGCCAGGAAATGTG  1980
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3046  TGTCACTGGAAGACAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTGTATGG  3105

Qy   1981  CCCCTTCTCTAAGAAAATGCCCTATTTGCAGGGGTATAATCAAGGGTATAGTCTTCGTACAT  2040
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3106  ACAAAGAAGTTTCTGTTGTATTTATTTTCCTTGTGGTCATCTGGTAGTAGTATGCCAGGAAATGTG  3165

Qy   2041  TTCTCTCTTAAAGAAAAATAGTTCTATATTTTTAAACCTGCATAAAAAAAGGTCTTTAAATAATT  2100
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3166  CCCCTTCTCTAAGAAAATGCCCTATTTGCAGGGGTATAATCAAGGGTATAGTCTTCGTACAT  3225

Qy   2101  GTTGAACACTTGAAGCCATCTAAAGTAAAAAAGGGAAATTATGAGTTTTTCAATTAGTAACA  2160
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3226  TTCTCTCTTAAAGAAAAATAGTTCTATATTTTTAAACCTGCATAAAAAAAGGTCTTTAAATAATT  3285

Qy   2161  TTCATGTTCTAGTCTGCTTTGGTACTAATAATCTTGTTTCTGAAAAGATGGTATCATATA  2220
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3286  GTTGAACACTTGAAGCCATCTAAAGTAAAAAAGGGAAATTATGAGTTTTTCAATTAGTAACA  3345

Db   3346  TTCATGTTCTAGTCTGCTTTGGTACTAATAATCTTGTTTCTGAAAAGATGGTATCATATA  3405
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Qy   2221  TTTAAATCTTAATCTGTTTATTTTACAAGGGAAGAGATTTANGTTTGGTGAACTATATATTAGTTAT  2280
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3406  TTTAAATCTTAATCTGTTTATTTTACAAGGGAAGAGATTTANGTTTGGTGAACTATATATTAGTTAT  3465

Qy   2281  GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTGTTATCTCCATCATTTCAGGAGTTACTGG  2340
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3466  GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTGTTATCTCCATCATTTCAGGAGTTACTGG  3525

Qy   2341  ATTTGTGTTGTTCTTTCAGAAAAGCTTTGAATACTAAATTATAGTGTTAGAAAAGAACTGGAAA  2400
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3526  ATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAATTATAGTGTTAGAAAAGAACTGGAAA  3585

Qy   2401  CCAGGAACTCTGGGAGTTCATCAGAGTTTATGGTCCGAATTGTCTTTTGCTTTCACTT  2460
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3586  CCAGGAACTCTGGGAGTTCATCAGAGTTTATGGTCCGAATTGTCTTTTGCTTTCACTT  3645

Qy   2461  GTGTTTAAAATAAGGAATTTTCTCTTATTTTCTCCCCCTAGTTTTGGTGAGAAACATCTCAA  2520
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3646  GTGTTTAAAATAAGGAATTTTCTCTTATTTTCTCCCCCTAGTTTTGGTGAGAAACATCTCAA  3705

Qy   2521  TAAAGTGCTTTAAAAATAAGGAAAAAAAAAAAAAAAA  2547
           |||||||||||||||||||||||||||
Db   3706  TAAAGTGCTTTAAAAATAAGGAAAAAAAAAAAA  3732
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Db   464 GTGTGAATGACAAGGTCAAAATGCTTCTGTTGTGTGGGCCTGATGTGGATAAACTGGAAACTAG 523

Qy   481 GAGACAGTCCTATTCAAAAGCATAAACACGTATATCCTAGCTGTAGCTTTATTCAGAATC 540
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   524 GAGACAGTCCTATTCAAAAGCATAAACACGTATATCCTAGCTGTAGCTTTATTCAGAATC 583

Qy   541 TGGTTTCAGCTAGTCTGGGATCCACCTCTTAAGAATACGTTCTCCAATGAGAACAGTTTTG 600
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   584 TGGTTTCAGCTAGTCTGGGATCCACCTCTTAAGAATACGTTCTCCAATGAGAACAGTTTTG 643

Qy   601 CACATTCATTATTCTCCCACCTTGGGAACATAGTAGCTTGTTGTCAGTGGTTCTTACTCCAGCC 660
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   644 CACATTCATTATTCTCCCACCTTGGGAACATAGTAGCTTGTTGTCAGTGGTTCTTACTCCAGCC 703

Qy   661 TTTCTCCCAAACCCTCTTAATTCAGATAGAGAGACAGTGTGAAGACANCTCTTCATCGGAGGACTAACC 720
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   704 TTTCTCCCAAACCCTCTTAATTCAGATAGAGAGACAGTGTGAAGACAGTGTGAAGACTAACC 763

Qy   721 CCTACAGTTATGCAAATGAGTACTGAAGAAGCCAGATTTCTTACCATATGTGGGCCAT 780
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   764 CCTACAGTTATGCAAATGAGTACTGAAGAAGCCAGATTTCTTACCATATGTGGGCCAT 823

Qy   781 TAACTTTTTTGTCACCATCAGAATTCAGAATTGGGCAAGAGAGCTGGTTTTAATATATAGGACCTGGAG 840
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   824 TAACTTTTTTGTCACCATCAGAATTGGGCAAGAGCTGGTTTTAATATATAGGACCTGGAG 883

Qy   841 ATAGGGTAGCCTGCTTTGCCTGTGGGAGCTCAGTACTCTGGGGAAACCAAAGGATGATG 900
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   884 ATAGGGTAGCCTGCTTTGCCTGTGGGAGCTCAGTACTCTGGGGAAACCAAAGGATGATG 943

Qy   901 CTATGTCAGAACACCGGAGGCCATTTCCCAACTGTCCATTTTTGGAAAAATTCTCTAGAAA 960
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Db   944 CTATGTCAGAAACACCGGAGGCCATTTCCCAACTGTCCATTTTTGGAAAAATTCTCTAGAAA 1003

Qy   961 CTCTGAGGTTTAGCATTTCAAATCTGACGATGCCACTTGACGATGAGGACGATGAGGAACAT 1020
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Db   1004 CTCTGAGGTTTAGCATTTCAAATCTGACGATGCCACTTGACGATGAGGAACAT 1063

Qy   1021 TTAAGTAACTGGCCATCTAGTTGTTCCAGTTCAGCGTTCAGCCTTGAGCGTCTGGTTTTT 1080
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1064 TTAAGTAACTGGCCATCTAGTTGTTCCAGTTCAGCGTTCAGCCTTGAGCGTCTGGTTTTT 1123

Qy   1081 ATTATGTGGGTCGCAAATGATGATGTCAAATGCTTTTGTTGTGATGGGTGGGCTTGAGGTGTT 1140
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1124 ATTATGTGGGTCGCAAATGATGATGTCAAATGCTTTTGTTGTGATGGGTGGGCTTGAGGTGTT 1183

Qy   1141 GGGAATCTGGAGATGAATCCATGGGTAGAACAATGCCAAGTCGTTTCCAAGGTGTGAGTTCT 1200
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1184 GGGAATCTGGAGATGAATCCATGGGTAGAACAATGCCAAGTCGTTTCCAAGGTGTGAGTTCT 1243

Qy   1201 TGATACGGAATGAAAGGCCCAAGAGAGTTTGTTGTTGATGAGATTCAGGTTGATATCCCTCATCTTC 1260
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Db   1244 TGATACGGAATGAAAGGCCCAAGAGTTTGTTGATGAGATTCAGGTTGATATCCCTCATCTTC 1303

Qy   1261 TTGAACAGCTGTGTGTCAACTCAGATACCACTGGAGAAGAAATGCTGACCCACCAATTA 1320
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Db   1304 TTGAACAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAATGCTGACCCACCAATTA 1363

Qy   1321 TTCATTTTTGGACCTGGAGAAGTTCTTCTTCAGATAGATGCTGTGTCATGATGAATACACCTGTGG 1380
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1364 TTCATTTTTGGACCTGGAGAAGTTCTTCTTCAGATAGATGCTGTGTCATGATGAATACACCTGTGG 1423

Qy   1381 TTAAATCTGCCTTGGAAATGGGGCTTTAATAGAGACCTGGGTGAAACAAAACAGTTCAAAGTA 1440
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1424 TTAAATCTGCCTTGGAAATGGGGCTTTAATAGAGACCTGGGTGAAACAAAACAGTTCAAAGTA 1483

Qy   1441 AAATCCTGACACTGGAGAGAACTATTAAAAACAGTTAATGATAATTGTTGTCAGCACTTCTTAA 1500
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Db   1484 AAATCCTGACACTGGAGAGAACTATAAAAACAGTTAATGATAATTGTTGTCAGCACTTCTTAA 1543

Qy   1501 ATGCTGAAGATGAAAAAGAGAGGAGGAGGAGAAAACACAGNCTGAAGAAGAAATGGCATCAG 1560
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Db   1544 ATGCTGAAGATGAAAAAGAGAGGAGGAGGAGGAAAAAACAGNCTGAAGAAGAAATGGCATCAG 1603

Qy   1561 ATGATTTGTCATTAATTCGGAAGAACACAGAATGGCTCTCTCTTTCAACAATTGACATGTGC 1620
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1604 ATGATTTGTCATTAATTCGGAAGAACACAGAATGGCTCTCTCTTTCAACAATTGACATGTGC 1663

Qy   1621 TTCCTTATCCTGGATAATCTTTTTTTAAAGGCCAATGTAATTAATAATAATAATAGGAACATGATATTA 1680
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Qy   1681 TTAAACAAAAAACACAGATACCTTTACAAGCCGAGAACTGATTGATACCCATTTTGGTTA 1740
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Db   1724 TTAAACAAAAAACACAGATACCTTTACAAGCCGAGAACTGATTGATACCCATTTTGGTTA 1783

Qy   1741 AAGGAAAATGCTGCGCGCCAACATCTTCAAAAACTGTCTAAAAGAAATTGACTCTACATTGT 1800
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Db   1784 AAGGAAAATGCTGCGCGCCAACATCTTCAAAAACTGTCTAAAAGAAATTGACTCTACATTGT 1843

Qy   1801 ATAAGAAACTTATTTGTGGATAAGAATATGAAGTATATCCAACAGAGAAGATGTTTCAGGTC 1860
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Db   1844 ATAAGAAACTTATTTGTGGATAAGAATATGAAGTATATCCAACAGAGAAGATGTTTCAGGTC 1903

Qy   1861 TGTCACTGGAAGACAAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTGTATGG 1920
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Db   1904 TGTCACTGGAAGACAAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTGTATGG 1963

Qy   1921 ACAAAGAAGTTTCTGTTGTATTTTAATTCCTTGTGGTCATCTGGTAGTATGCCAGGAAGTG 1980
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Qy   1981 CCCCTTCTCTAAGAAAATGCCCTATTTGCAGGGGTATAAATCAAGGGTATACTGTTCGTACAT 2040
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2024 CCCCTTCTCTAAGAAAATGCCCTATTTGCAGGGGTATAAATCAAGGGTATACTGTTCGTACAT 2083

Qy   2041 TTCTCTCTTAAAGAAAAAAATAGTCTATATTTTAAACCTGCATAAAAAAAGGTCTTTAAAATATT 2100
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Db   2084 TTCTCTCTTAAAGAAAAAAATAGTCTATATTTTAAACCTGCATAAAAAAAGGTCTTTAAAATATT 2143

Qy   2101 ACAAAGAACACTTGAAGCCATCTAAAGTAAAGTAAAAAGGGAATTATGAGTTTTCAGGTAGTATGTTAACA 2160
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Db   2144 GTTGAACACTTGAAGCCATCTAAAGTAAAGTTGTTAGGTGCATCATTCAAGGAGTTACTGG 2203

Qy   2161 TTCATGTTCTAGTCTGCTTGGTACTTAATAATCTTGTTTCTGAAAAGAATGGTATCATATA 2220
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Db   2204 TTCATGTTCTAGTCTGCTTGGTACTTAATAATCTTGTTTCTGAAAAGAATGGTATCATATTA 2263

Qy   2221 TTTAAATCTTAAATCTGTTTAATTACAAAGGAAGAGATTTATGTTTGGTGAACTAATTAGTTAT 2280
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Db   2264 TTTAAATCTTAAATCTGTTTAATTACAAAGGAAGAGATTTATGTTTCGGTGAACTATAATATAT 2323

Qy   2281 GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTGTTATTGGTCATATTTCAGGGATTACTGG 2340
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Db   2324 GTATGTGTACCTAAGGGAGTAGCGTCCNCTGCTTGTTGTTATCATTTCAGGAGTTACTGG 2383

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Db   2444 CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCGAAATTGTCTTTTGGTGCTTTTCACTT 2503

Qy   2461 GTGTTTAAATAAGGATTTTTTCTCTTATTTCTCCCCCTAGTGTTGTTGTGAAACATCTCAA 2520
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Db   2504 GTGTTTTAAAATAAGGATTTTTCTCTTATTTCTCCCCCTAGTGTTTGTTGTGAAACATCTCAA 2563

Qy   2521 TAAAGTGCTTTAAAAA 2536
          |||||||||||||
Db   2564 TAAAGTGCTTTAAAAA 2579

Patent No. 6541457

GENERAL INFORMATION:

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APPLICANT: MacKenzie, Alexander E.
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APPLICANT: Liston, Peter

TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
TITLE OF INVENTION: PROBES, AND DETECTION METHODS

FILE REFERENCE: 07891/003003

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EARLIER FILING DATE: 1996-08-05
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EARLIER FILING DATE: 1995-12-22
EARLIER APPLICATION NUMBER: 08/511,485
EARLIER FILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 45
SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 7
LENGTH: 2580
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: variation
LOCATION: (2412)...(2412)
OTHER INFORMATION: N may be any nucleotide

US-09-201-936-7

Query Match   97.1%; Score 2514.8; DB 4; Length 2580;
Best Local Similarity   99.4%; Pred. No. 0;
Matches 2522; Conservative 0; Mismatches 14; Indels 0; Gaps 0;

Qy   1 TCTAAGTAGTATCTTGGGAAATTCAGAGAGATACTCATCATCCTACCTGAATATAAACTGAGAT 60
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Db   364 TCCCCGCCGCGGGGTGCCTGTCTCTCAGAAAGGAGAGTCTTGCTTGCTGTTGCTGTTTATTATACTG 423

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Qy   481 GAGACAGTCCTATTCAAAAGCATAAACAGCTATAATCCTAGCTGTAGCTTTTTATTCAGAATC 540
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy   541 TGGTTTCAGCTAGTCTGGGATCCACCTCTAAGAATACGTCTCCAATGAGAAACAGTTTTG 600
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Db   601 CACATTCATTATATCTCCCACCTTGGAACATAGTAGCTTGGTTGTTCAGTGGTTCTTACTCCAGCC 660

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Db   644 CACATTCATTATATCTCCCACCTTGGAACATAGTAGCTTGGTTGTTCAGTGGTTCTTACTCCAGCC 703

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Db   764 CCTACAGTTATGCAAATGAGTACTGAAGAAGCCAGATTTCTTACCTTACCCATATGTGGCCAT 823

Qy   781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGACGTGGTTTTTTTATTATATAGGAACTAAC 840
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Db   824 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGACGTGGTTTTTTTATTATATAGGAACTAAC 883

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Db   944 CTATGTCAGAACACCCGGAGGGCAATTTTCCCAAACTGTCCCCATTTTTGGAAAAATTCTCTAGAAA 1003

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Qy 1081 ATTAGTGTGGGGTCGCAAATGAATGATGATGAGTCAAATGCTTTTGTTCTTGAATGGTCGGCTTGGTCGGCTTTGAAGGTGTT 1140
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Db 1124 ATTATGTGGGGTCGCAAATGAATGATGAGTCAAATGCTTTTGTTCTTGATGGTCGGCTTGGTTGGTGTT 1183

Qy 1141 GGGAATCTGGAGATGACCCATGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGGAGTTCTC 1200
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Db 1184 GGGAATCTGGAGATGACCCATGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGGAGTTCT 1243

Qy 1201 TGATACGGAATGAAAGGCCAAGAGTTTGTTGTTGATGAGATTCAAGGTAGTAGTATCCTCATCTTC 1260
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Db 1244 TGATACGGAATGAAAGGCCAAGAGTTTGTTGTTGATGAGATTCAAGGTAGTATTCTCATCTTC 1303

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Db 1304 TTGAACAGCTGTTGTCAACTTCAGATACCACTGGGAGAAGAAAATGCTGACCCCACCAATTA 1363

Qy 1321 TTCATTTTGGACCTGGAGAAGAAGTTCTTCAGAAGATGCTGTCGTCATGATGAATAACACCTGTGG 1380
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1364 TTCATTTTGGACCTGGAGAAGAAGTTCTTCAGATGATCATGATGAATAACACCTGTGG 1423

Qy 1381 TTAAAATCTGCCCTGGGAAATGGGCTTTAATAGGAACAAAACAGTTCAAAGTA 1440
         ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1424 TTAAAATCTGCCCTGGGAAATGGGCTTTAATAGGAACAAAACAGTTCAAAGTA 1483

Qy 1441 AAATCCTGACAACTGGAGAGAACTATAAAACAGTTAATGAATGTGTCAGCACTTCTTA 1500
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1484 AAATCCTGACAACTGGAGAGAACTATAAAACAGTTAATGAATGTGTCAGCACTTCTTA 1543

Qy 1501 ATGCTGAAGATGAAAAAAGAGAGGAGGAGAAAAGTGAAGAAAATGGCATCAG 1560
         ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1544 ATGCTGAAGATGAAAAAAGAGAAGAGAGGAGGAGAAAAGTGAAGAAAATGGCATCAG 1603

Qy 1561 ATGATTTGTCATTAATTCCGGAAGAACAGAGAATGGGCTCTCTTTCAAACAATTGTGC 1620
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1604 ATGATTTGTCATTAATTCGGAAGAACAGAGAATGGGCTCCTTTCAAACAATGACATGTGC 1663

Qy 1621 TTCCTATCCTGGATAAATCTTTTAAAGGCCAATGTAATTAATAAAACAGGAAACATGATATTA 1680
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 1664 TTCCTATCCTGGATAAATCTTTTAAAGGCCAATGTAATTAATAAAACAGGAAACATGATATTA 1723

RESULT 9

PCT-US95-05922A-1

; Sequence 1, Application PCT/US9505922A
; GENERAL INFORMATION:
;   APPLICANT: HE, ET AL.
;   TITLE OF INVENTION: Human Inhibitor of Apoptosis Gene 1
;   NUMBER OF SEQUENCES: 8
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
;   CECCHI, STEWART & OLSTEIN
;   ADDRESS:
;   STREET: 6 BECKER FARM ROAD
;   CITY: ROSELAND
;   STATE: NEW JERSEY
;   COUNTRY: USA
;   ZIP: 07068
;   COMPUTER READABLE FORM:
;   MEDIUM TYPE: 3.5 INCH DISKETTE
;   COMPUTER: IBM PS/2
;   OPERATING SYSTEM: MS-DOS
;   SOFTWARE: WORD PERFECT 5.1
;   CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: PCT/US95/05922A
;   FILING DATE: 11 MAY 1995
;   CLASSIFICATION:
;   PRIOR APPLICATION DATA:
;   APPLICATION NUMBER:
;   FILING DATE:
;   ATTORNEY/AGENT INFORMATION:
;   NAME: FERRARO, GREGORY D.
;   REGISTRATION NUMBER: 36,134
;   REFERENCE/DOCKET NUMBER: 325800-292
;   TELECOMMUNICATION INFORMATION:
;   TELEPHONE: 201-994-1700
;   TELEFAX: 201-994-1744
; INFORMATION FOR SEQ ID NO: 1:
;   SEQUENCE CHARACTERISTICS:
;   LENGTH: 1435 BASE PAIRS
;   TYPE: NUCLEIC ACID
;   STRANDEDNESS: SINGLE
;   TOPOLOGY: LINEAR
;   MOLECULE TYPE: CDNA
; PCT-US95-05922A-1

Query Match 55.2%;   Score 1430.2;   DB 5;   Length 1435;
Best Local Similarity 99.8%;   Pred. No. 8.9e-297;
Matches 1432; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy   726 AGTTATGCAATGAGTAGTACTGAAGAGAGCCAGAGATTTCTTACCTACCATATGTGGCCATTAACT 785
           |||||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||
Db   1 AGTTATGCAATGAGTACTGAAGAGAGCCAGAGATTTCTTACCTACCATATGTGGCCATTAACC   60

Qy   786 TTTTTGTCACCATCAGAATTGGCAAGAGCTGGGTTTTTTATTATATATAGGACCCTGGAGGATAGG 845
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 TTTTTGTCACCATCAGAATTGGCAAGAGCTGGGTTTTTTATTATATATAGGACCCTGGAGGATAGG 120

Qy   846 GTAGCCTGCTTTGCCTGTGGGTGGGGAAGCTCAGTAACTGGTCAGTAACTGGGAA   905
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121 GTAGCCTGCTTTGCCTGTGGGTGGGGAAGCTCAGTAACTGGTCAGTAACTGGGAA   180

Qy   906 TCAGAACACCCGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAATTCTCTAGAAACTCTG   965
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181 TCAGAACACCCGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAATTCTCTAGAAACTCTG   240

Qy   966 AGGTTTAGCATTTCAAAATCTGAGCAATGCAGACACAATGCAGCTCGAATGAGAACATTTATG   1025
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241 AGGTTTAGCATTTCAAAATCTGAGCAATGCAGACACAATGCAGCTCGAATGAGAACATTTATG   300

Qy   1026 TACTGGGCCATCTAGTGTGTTCCAGTTCAGCCTGAGCCTTGCAAGTGCTGGTTTTTTATTTAT   1085
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   301 TACTGGGCCATCTAGTGTGTTCCAGTTCAGCCTGAGCCTTGCAAGTGCTGGTTTTTTATTTAT   360

Qy   1086 GTGGGTCGCAATGATGATGATCAAATGCTTTTGTTGTTGTGATGGTGGCCTTGAGGTGTTGGGAA   1145
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   361 GTGGGTCGCAATGATGATGATCAAATGCTTTTGTTGTTGTGATGGTGGCCTTGAGGTGTTGGGAA   420

Qy   1146 TCTGGAGGATGGATCCATGGGTAGAACATGCCAAGGTGGTTTCCAAGGTGTGAGTTCTTGATA   1205
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   421 TCTGGAGGATGGATCCATGGGTAGAACATGCCAAGGTGGTTTCCAAGGTGTGAGTTCTTGATA   480

Qy   1206 CGAATGAAAGGCCAAGAGTTTGTTGATGAGGATTCAAGGTAGATATCCTCATCTTCTTCATTT   1265
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   481 CGAATGAAAGGCCAAGAGTTTGTTGATGAGGATTCAAGGTAGATATCCTCATCTTCTTCATTT   540

Qy   1266 CAGCTGTTGTCAACTTCAGATACCACTGGAGGAAGAAAAATGCTGACCCACCAATTATTCAT   1325

Qy   1681 TTAAACAAAAAACACAGATACCTTTACAGAAGCGGAGGAGGAACTGAGTTGGATTCGATACCATTTGGTTA   1740
Db   1724 TTAAACAAAAAACACAGATACCTTTACAGAAGCGGAGGAGGAACTGAGTTGGATTCGATACCATTTGGTTA   1783

Qy   1741 AAGGAAAATGCTGCGGCCAACATCTTCAAAAAAACTGTCTAATAGAAATTGACTTCACATTGT   1800
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1784 AAGGAAAATGCTGCGGCCAACATCTTCAAAAAAACTGTCTAATAGAAATTGACTTCACATTGT   1843

Qy   1801 ATAAGAACTTATTTGTGGATAAGAAATATGAAGTGATATTCCAACAGAAGATAGTTTCAGGTC   1860
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1844 ATAAGAACTTATTTGTGGATAAGAAATATGAAGTGATATTCCAACAGAAGATAGTTTCAGGTC   1903

Qy   1861 TGTCACTGGAAGAACAAATTGAGGAGGGTTGCAAGAAGACCAACTTGTAAAGTGTGTATGG   1920
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1904 TGTCACTGGAAGAACAAATTGAGGAGGGTTGCAAGAAGACCAACTTGTAAAGTGTGTATGG   1963

Qy   1921 ACAAAGAGAGTTTCTGTTGTATTAATTCCTTGTGGGTCATCTGGTAGTATGCCAGGAATGTG   1980
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1964 ACAAAGAGAGTTTCTGTTGTATTAATTCCTTGTGGGTCATCTGGTAGTATGCCAGGAATGTG   2023

Qy   1981 CCCCTTCTCTAAGAAAAATGCCCTATTTGCAGGGGTATAAATCAAGGGTACTGTTCGTACAT   2040
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2024 CCCCTTCTCTAAGAAAAATGCCCTATTTGCAGGGGTATAAATCAAGGGTACTGTTCGTACAT   2083

Qy   2041 TTCTCTCTTAAAGAAAAAATAGTCTATATATTTTAACCTGCATAAAAAGGTCTTTAAAAATATT   2100
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2084 TTCTCTCTTAAAGAAAAAATAGTCTATATATTTTAACCTGCATAAAAAGGTCTTTAAAAATATT   2143

Qy   2101 GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGAAATTATGAGTTTTCAAATTAGTAGTAACA   2160
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2144 GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGAAATTATGAGTTTTCAAATTAGTAGTAACA   2203

Qy   2161 TTCATGTTCTAGTCTTGGTACTAATAATCTTGTTTCTGAAAAGATGGTATCATCATATA   2220
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2204 TTCATGTTCTAGTCTGCTTTGGTACTAATAATCTTGTTTCTGAAAAGATGGTATCATCATATA   2263

Qy   2221 TTTTAATCTTAATCTGTTTATTTACAAGGGACAGATTTATGTTTTGGTGAACTATATTAGTTAT   2280
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2264 TTTTAATCTTAATCTGTTTATTTACAAGGGACAGATTTATGTTTTGGTGAACTATATTAGTTAT   2323

Qy   2281 GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTATGCNTCANTTCAGGAGTTACTGG   2340
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2324 GTATGTGTACCTAAGGGAGTAGCGTCNCTGCCTTGTTATGCNTCANTTTCAGGAGTTACTGG   2383

Qy   2341 ATTTGTTGTTCTTTCAGAAAGCTTTGAAAACTAAATTATAGTGTAGAAAGAAGAACTGGGAAA   2400
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2384 ATTTGTTGTTCTTTCAGAAAGCTTTGAAAACTAAATTATAGTGTAGAAAGAAGAACTGGGAAA   2443

Qy   2401 CCAGGAACTCTGGAGTTCATCAGAGTTGTCCGAATTGTCCTTTTGGGTGGCTTTTCACTT   2460
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2444 CCAGGAACTCTGGAGTTCATCAGAGTTGTCCGAATTGTCCTTTTGGGTGGCTTTTCACTT   2503

Qy   2461 GTGTTTAAAATAAGGATTTTTTCTCTTATTTCTCCCCTAGTTGTTTGTTGTGAGAAACATCTCAA   2520
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2504 GTGTTTAAAATAAGGATTTTTTCTCTTATTTCTCCCCTAGTTGTTTGTTGTGAGAAACATCTCAA   2563

Qy   2521 TAAAGTGCTTTAAAAA   2536
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Db   2564 TAAAGTGCTTTAAAAA   2579

RESULT 10
US-08-569-749-13
Sequence 13, Application US/0856974
Patent No. 6187557

GENERAL INFORMATION:

; APPLICANT:
; APPLICANT: Rothe, Mike
; APPLICANT: Goeddel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco

; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749
; FILING DATE:
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Brezner, David J.
; REGISTRATION NUMBER: 24,774
; REFERENCE/DOCKET NUMBER: A-62464/DJB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415)781-1989
; TELEFAX: (415)398-3249
; INFORMATION FOR SEQ ID NO: 13:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2862 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA

US-08-569-749-13

Query Match   52.0%;   Score 1345.2;   DB 3;   Length 2862;
Best Local Similarity 76.6%; Pred. No. 1.6e-278; Indels 79; Gaps 13;
Matches 1844; Conservative 0; Mismatches 483;

Qy 140 TGTGAAGAAAATTTCATGTGAATCTTTTAGCTATCAAACAGCACTGTCACCTACTCATGCA 199
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Db 420 TGGTGAGAGAACTTCATCTGGAAGTTAAGCGGTCAGAAAATACTATT--ACTACTCATGGA 477

Qy 200 CAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTCGTATCAAAACATTAAGAGTAATAAT 259
        ||||| |||||| ||||| ||| |||||||||||| || ||||| |||| |||||| ||
Db 478 CAAAACTGTCTCCCAGGACTCGGCCACCAAGGTACCTTACACCAAAAACTTAAACGTATAAT 537

Qy 260 GGAAGATAGCACGATCTTGTCAGATTGGACAAAACAGCACAAAAAAAATGAAGTATCA 319
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Db 538 GGAGAAGAGCACAAATCTTGTTCAAATTGGACACAAAGGAAGAAAAAATGAAAGTTTGA 597

Qy 320 CTTTTCCTGTGAACTCTACAGAATGTCTACATAATTCCCGCGCCGGGGTGCCTGT 379
        |||||||||||||| || || | |||| | | ||||||||| |||||||||||||
Db 598 CTTTTCGTGTGAACTCTACCGAAATGTCTACATATTTCCCAGGGGGAGTTCCTGT 657

Qy 380 CTCAGAAAAGGAGTCTTTGCTCGTCGTCTGGTTTTTTATTATTATTGACAAGTCAA 439
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Db 658 CTCAGAGAGGAGTCTGGGCTGGCTCCTTTTATTATACAGGTGTGAATGACAAAAGTCAA 717

Qy 440 ATGCTTCTGTTGTGGCCTGGATAACTGGAAACTAGGAGAGACAGTCCTATTCAAAA 499
        ||| |||||| | |||||||| ||| |||||| ||||||||||||||| ||||||
Db 718 GTGCTTCTGCTGTGGCCCTGATGTTGGAATAACTGGAAACAAAGGGGACAGTCTGTGAAAA 777

Qy 500 GCATAAACAGCTATATCCTAGCTGTAGCTTTATTCAGAATCTGGTTTCAGCTAGTCTGGG 559
        || |||| ||||| | |||| ||| ||||||||||||| ||||| ||||||||||||||||
Db 778 GCACAGACAGTTCTATCCCAGCTGCAGCTTTGTACAGACTCGCTTCAGCCAGTCTGCA 837

Qy 560 ATCCACCTCTAAGAATAGTCTCCCAATGAGAGAAACAGTTTTGCACATTCATTATCTCCCAC 619
        | | ||||| ||||| ||| ||||| ||| |||||||||||||| || ||||||| ||||
Db 838 GTCTCCATCTAAGAATATGTCTCCTGTGAAAAAGTAGATTTGCACATTCGTCACTCTGGA 897

Qy 620 CTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC 679
        |||| |||||| |||||| ||||||| || |||||||||||
Db 898 ACGAG------------GTGGGCATTCACTCCAACCTGTGTCTTAGCTGCCCTTTAA 939

Qy 680 TTCTCAGAGCAGTTGAAGCACATCTTTCATCGAGGACTAACCCCTACAGATTTATTGCAATCAG 739
        ||||| |||||| ||||| | ||| ||| |||||| |||||||||||||||| |||||||||||
Db 940 TTCTCAGACAGTGGAGGAGACTTC---TCATCAAGGATGGATCCCTGCAGCCCATCTGCGCTATGGTAG 996

Qy 740 TACTGAAGAAGCCAGATTTCTTCACTACCATATGTGGGCCATTAAACTTTTTTGTCACCATC 799

Qy   1814 TGTGGATAAGAATATGAAGTGTATTCCAACAGAAGAAGACGTTTCAGGCTTGTCATTGGAAGA 1873
           ||||| |||||||
Db   2014 AGCCAACATCTTCAAAAAACTCTCTGAAGGAAATTGACTCCA-GTTATATGAAAAACTTATT 2073

Qy   1754 GGCCACACATCTTCAAAAAACTGTCTCTAAAAGAAAATTGACTCTAGAATTGTATAAGGAAGTTTC 1813
           ||||||||| |||||
Db   1954 ACAGATACCCTTACAAGCGAGAGAACTGATTGATACCATATTTGGTTAAAGGAAATGCTGC 2013

Qy   1694 ACAGATACCTTTACAAGCGAGAGAACTGATTGATACCATATTTGGTTAAAGGAAATGCTGC 1753
           ||||||||| |||||
Db   1894 TAATTCTTCTTGAGGGCCAGTGTAATTACAAAACAGAGAACATGATATTAATTAAACAAAAAAC 1953

Qy   1634 TAATTCTTTTAAGGCCCAAATGTAATTACAAAACAGAGAACATGATATTAATTAAACAAAAAAC 1693
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Db   1834 GATTCGGAAGAATAGAAATGGCCCTCTTCAAACAGTTGACACAATGTCCTTCCTATCCTGGA 1893

Qy   1574 AATTCGGAAGAACAGAATGGCTCTCTTTCAAACAATTGACAATGTGTGCTTCCTATCCTGGA 1633
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Db   1774 GAGAGAAGAGGAGGAAAAGACAGAGACTGAAGAGATGCCATCAGGTGACTTATCACT 1833

Qy   1514 AAAAAGAGGAGGAGAGGAGGAGGAAAAACAAGCTGAAGAAATGCCATCAGAGATGAGATTTGTCATT 1573
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Db   1714 TGGTGAGAACTACAGGAGACCGTCAATGATAATGTCTCAGTACTTTTGAAATGCTGAAAGATGA 1773

Qy   1454 TGGAGAGAACTATAAAAACAGTTAATGATATATGTGTCAGCACTTCTAAAATGCTGAAGATGA 1513
           ||||||||| |||||
Db   1654 GGAAAATGGGCTTCAGTGAGCGCCTGGGTTGACGACGCCAGATCCTGGCCAC 1713

Qy   1394 GGAAAATGGGCTTTAATAGAGAGAACCTGGTGAAACAACAGTTCAAAGTAAAAATCCTGACAAC 1453
           ||||||||| |||||
Db   1597 TGGAGAAAG---TTCGGAAGATGTCGCTCATCATGAGTGCACGC-CTCTGGTTTTAAAGCAGCCTT 1653

Qy   1334 TGGAGAAAGTTCTTCAGAAGGATGCTGTGTCATGTGGGTTTAAAATCTGCCTT 1393
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Db   1537 TTCAGACACCCCAGGAGAGAAAAATGCTGACCCTACAGAGTCAGTGGTGCATTTTGGGCCC 1596

Qy   1280 TTCAGATACCACTGGGAGAGAGAGAAAAAATGCTGACCC-----ACCAATTATTCATTTGGACC 1333
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Db   1477 GGAGTTTGTTGATGAGATTCAAGCTAGAGATATCCTCATCTTCTTGAGCAGCTGTTGTCCAC 1536

Qy   1220 AGAGTTTGTTGATGAGATTCAAGGTAGAGATATCCTCATCCTTCTTGAACAGCTGTGTTGTCAAC 1279
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Db   1417 CTGGATAGAACACGCCCAAAATGGTTTCCAAGGTGTGAGTGTTCTTTGATACCGGATGAAGGGTCA 1476

Qy   1160 ATGGGTAGAACATGCCCAAGGTGGTTTCCAAGGTGTTGATCAGAATGAAATGAAAGGCCCA 1219
           ||||||||| |||||
Db   1357 TGAGTGTCAAGTGCTTTTGTTTGTTGTGATGGGTGGGTGTTGGGAATGTGGGTGTGAGGGCCTTGA 1416

Qy   1100 TGATGTCAAAATGCTTTGTTGTTTGTGATGGGTGGTGTTGGGAATGTGGGAATGATCC 1159
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Db   1297 TGTTCCCTGTTCAGCCCGAGCAGTTGCAAGTGCTGGAATCTATTACGTTGGATCGCAATGA 1356

Qy   1040 TGTTCCAGTTCAGCCTGACGACGGCTTGCAAGTGCTGGTTTTTATTATGTGGGTCGGCAATGA 1099
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Db   1237 AAATCTAAGTATGCAGACACACTCGCCTCGCAGTGGCCACCTAG 1296

Qy   980 AAAATCTGAGCATGCAGACAACATGCCAGCTCGGAATGAGAACATTTATGTACTGGCCATCTAG 1039
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Db   1177 ACATTTTCCCACTGTCCAATTTCTGGAAAAATACTTCAGAAAAACACAGAGGTTTAGTATATC 1236

Qy   920 GCATTTTCCCAACTGTCCATTTTTTGGAAAAATTCTCTAGAAACTCTGAGGTTTAGCATTTC 979
           ||||||||| |||||
Db   1117 CTGTGGGTGGGAAACTGAGCAACTGGGGAACCAAAGGATGATGCTATGTGTCAGGCACCGCAG 1176

Qy   860 CTGTGGTGGGGAAGTCAGTAACTGGGGAAACCAAAGGATGAT GCTAtGTCAGAaCACCGGAG 919
           ||||||||| |||||
Db   1057 AGAGCTGGCCAGAGCTGGCTTCTATTACATAGGGCCTTGTTTTGC 1116

Qy   800 AGAATTGGCAAAGAGCTGGTTGTTTTATTATATAGGGGTAGGCCTTGGCCTTGC 859
           ||||||||| |||||
Db   997 TACAGAAGAGAGGGCCAGATTTCTTTACTTACGTATGTGGGGCTTTAAGTTTTCTGTCACCAGC 1056

...sequence alignment continues...

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CLASSIFICATION:
  PRIOR APPLICATION DATA:
    APPLICATION NUMBER: U.S. Serial Nos. 08/512,946 & 08/569,749
    CLASSIFICATION:
  ATTORNEY/AGENT INFORMATION:
    NAME: Brezner, David J.
    REGISTRATION NUMBER: 24,774
    REFERENCE/DOCKET NUMBER: A-62464/DJB
  TELECOMMUNICATION INFORMATION:
    TELEPHONE: (415)781-1989
    TELEFAX: (415)398-3249
INFORMATION FOR SEQ ID NO: 13:
  SEQUENCE CHARACTERISTICS:
    LENGTH: 2862 base pairs
    TYPE: nucleic acid
    STRANDEDNESS: single
    TOPOLOGY: linear
    MOLECULE TYPE: cDNA
PCT-US96-12860-13

Query Match    52.0%;  Score 1345.2;  DB 5;  Length 2862;
Best Local Similarity 76.6%;  Pred. No. 1.6e-278;
Matches 1844;  Conservative 0;  Mismatches 483;  Indels 79;  Gaps 13;
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Qy   140 TGTGAACAAATTTCATGTGAAATGTTTTAGCTATCAAACAGCACTGTCACCTACTCATGCA 199
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Db   420 TGGTGAGAAACTTCATCTGGAAGTTTAAGGGGTCAGAAATACTATT--ACTACTCATGGA 477

Qy   200 CAAAACTGCCTCCCCAAAGACTTTTTCCCAGGTCCCTCGTATCAAAAACATTAAGAGTATAAT 259
           ||||||||||||| ||||||| ||||||| ||||||||||||||||||||||||||
Db   478 CAAAACTGTCTCCCCAGGACTCGCGCCAGGTACCTTACACCAAAAACTTAAACGTTATAAT 537

Qy   260 GGAAGATAGCACGATCTTGTTCAGATTGGACAAACAGCAACAAAACAAAAAAAATGAAAGTATGA 319
           ||||| |||||||| ||||||||||||||||| |||||||| |||||||||||||||||
Db   538 GGGAGAAGAGACACAATCTTGTCAAATTGGACAGGAGAGGCGAAGAAAAATGAAGTTTGA 597

Qy   320 CTTTTCCTGTGAACTCTACAGAAATGTTCTACAGATATCAAACTTTCCCCGGCCGGGGGGGTGCCTGT 379
           |||||||| |||||||||||||||||||||||||||||| |||||||||||||| |||||||
Db   598 CTTTTCGTGTGAACTCTTCACCGAAGTCTTCTACATAATTCAGCTTTTCTTTGAAAA 657

Qy   380 CTCAGAAAAGGAGTCTTGCTCGTGCTCGGTTTTTTTTATTTATAACTCGGTTGGAATGACAAGGTCA 439
           |||||||| |||||||||||||||| |||||| |||| |||||||||||||
Db   658 CTCAGAGGAGGAGTCTCGGCTGTCGTCGTGCTTTTATTTATTTATACAGGTGTGAATGACAAAGTCAA 717

Qy   440 ATGCTTCTGTTGTTGGGCCCTGATCGTCCGATAACTGGAGAACTGGACAGTCCTATTCACAAA 499
           ||||||| || |||||||||||||||| |||||||||||||||||||||||||||||||||
Db   718 GTGCTTCTGCTGTGGCCCTGATGTTGGATAACTGGAAAACATGGGGAACAGTCCTGTTGAAAA 777

Qy   500 GCATAAACAGCTATATCCTGGTGTAGCTTATTCAGAA TCTGGGTTTCAGCTAGTCTGGG 559
           |||||||||| ||||||| |||||||| ||||||||| ||||||||||||||||||||||
Db   778 GCACAGACAGTTCTATCCCAGCTGCAGCTTTGTACAGAC CTGCTTTCAGCCCAGTCTCA 837

Qy   560 ATCCACCTCTAAGAAATAAGGTCTCCAAATGAGAGAAACAGTTTGGCACATTCATTATCTCCCAC 619
           |||||| |||||||||| ||||||| ||||| |||||||||| ||||||||||||||||||
Db   838 GTCTCCATCTAAGAAATATGTCTCCTGTGTGAAAAAGTATGGCACATTCGTCACCTCTCGGA 897

Qy   620 CTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTCCCAGCCCTTCTCCCAAACCCTCTTAA 679
           ||||||| ||||| ||| ||||||||||||||||||||||||||||||||||||||||||||
Db   898 ACGAG--------------GTGGCAATTCACTCCCAAC CTGTGTCCTTCTCCTAGCCCTCTTAA 939

Qy   680 TTCTAGAGCAGTTGAAGACATCTTCTTACCTACCACATATGTGGGCCACATTAACTTTTTTTGTCACCATC 739
           |||||||||||||||||| || |||||||||| ||||||||||||||||||||||||||||||
Db   940 TTCTRGAGCAGTGGAAGACTTC---TCATCAAGGATGGGAACCCTGCAGCTAGCCATGAG 996

Qy   740 TACTGAAGAAGCAGCGAGATTTCTTACCTACCACCATATGTGGGCCA... 799
           |||||||||||||||||||||||||||||||||||||||||||||||||
Db   997 TACAGAAGGAGGCCAGATTCCTTACTTACAGTATTTCAGGCTTGCATTGGGAAGA 1056

Qy   800 AGAATTGGCAAGAGCTGGTTTTTTATTATATAGGACCTGGAGATAGGGTAGCCTGCTTTGC 859
           ||||||||||||||||| ||||| ||| |||| |||||||||||||||||| |||||||||
Db   1057 AGAGCTGGCCCAGAGCTGGCTTCTTATTACATAGGGCCTGGCAGCAGGGGTGCCCTGGTTTC 1116

Qy   860 CTGTGGTGGGAAGCTCAGTAACTGGGACCAAAGAATGAGGATGATGCTATGTGTCAGAACACACCCGAG 919
           |||| |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1117 CTGTGGTGGGAAAACTGAGCAACTGGGAGGAACCAAAGGATGATGCTATGTGTACTGGCCATCGA 1176

Qy   1160 ATGGGTAGGAACAATGCCAAGTGCTTCCAAGGTGTGAGTGCTGCTTGATAAAGCCCA 1219
Db   1417 CTGGATAGAACACGCCAAATGGTTTCCAAGGGTGTGAGTTCTTGAATACGGATCAAGGTCA 1476

Qy   1220 AGAGTTTGTTGATGAAGATTCAAGGTAGAGTATCCTCATCTTCTCTGAACAGCTGTTGTCAAC 1279
Db   1477 GGAGTTTGTTGATGAGATTCAAGCTAGAGATAATCCTCATCTTCTTGAGCAGCTGTTGGCAC 1536

Qy   1280 TTCAGATACCAACTGGAGGAAGAAAAATGCTGACCC---... 1333
Db   1537 TTCAGACACCCCCAGGAGAGAAAAATGCTGAAGCCCTTACAGAGACACAGTGGCATTTTGCCC 1596

Qy   1334 TGGGAAAGTTCTTCAGAAGATGCTGTCATGATGAATCACACCTGTTGTAAAATCTGCCTT 1393
Db   1597 TGGGAGAAAG---TTCGGAAGATTCGTCATGATGAGCACGGCCCTGGTTAAAGCAGCTT 1653

Db   1117 |||||||| |||||||||||||||||||||||||||||||||||||||| 1176
Db   920 GCATTTTCCCAAACTGTGCCATTTTTGGAAAATTCTCTAGAAAACTCGAGGGTTTAGCATTTC 979

Db   1177 ACATTTTCCCACTGTCCATTTCTGGAAAATACTTCAGAAAACACAGAGGTTTAGTATATC 1236

Db   980 AAAATCTGAGCATGCAGACAACATGCAGCTCGAATGAGAGAACATTATTGTACTGGCCATCTAG 1039
Db   1237 AAATCTAAGATATGCAGACAACACTCCTGCTCGGATGAGACAGTTGGGTTGGGAATCGATCC 1296

Db   1040 TGTTCCCAGTTCAGCCTGAGCAGCTTGCAAGTGCTTGGTTTTTATTATGTGGGTTCGCAATGA 1099
Db   1297 TGTTCCTGTTCAGCCCGAGCAGCTTGCAAGTGCTGGAATCGCAATGA 1356

Db   1100 TGATGTCAAATGCTTTGTGTTGTGGATGGTGGGATCTGGAGAAATCTGCCTT 1159
Db   1357 TGATGTCAAGTGCTTTTGTTTTGTTTGGATGGTGGGGTTGGGAATCTGGAGGATGATGACC 1416

Qy   1394 GGAAATGGGCTTTAATTAGAGACACCTGTGTGAAACACAGTAAAAATCCTGACAAAC 1453
Db   1654 GGAAATGGGCTTCAGTTAGGGACCCTGGTGAGACAACGGGTTCAGCGACCTGGCCAC 1713

Qy   1454 TGGAGAGACTATAAAAACAGTTAATGATATTGTGTCAGCACTTCTAAAATGCTGAAGATGA 1513
Db   1714 TGGTGAGAACTACAGGACCCGTCAATGATAATTGTCTCAGTGACTTTTGAAAGCTGAAGATGA 1773

Qy   1514 AAAAAGAGAGGAGGGAAGGAAAAAACAAGCTGAAGAAAATGGCATCAGATGAGATTTTGTCATT 1573
Db   1774 GAGRAGGAGGAGGAAAGGAAAGACAGAACTGGCATCCAGGTGACTTATCACT 1833

Qy   1574 AATTCGGGACAGAAACAGAATGGCTCTCTCTTTCAACAATTGACATGTGGTGCTTCCTATCCTGGA 1633
Db   1834 GATTCGGGAAGAATAGAATAGGCCCTCTTCAACAGTTGACACATGTTCCTTCCTATCCTGGA 1893

Qy   1634 TTAATCTTTTAAAGGCCAATGTAATTAATAAACAGGACAATGATATTGTATTATTAAACAAAAAC 1693
Db   1894 TAATCTTTCTTTGGAGGCCAGTGTAATTACAAAACAATGATATTTAGACAGAAAC 1953

Qy   1694 ACAGATACCTTTACAAGGCGAGAGGAACTGATTGGATACCATTTGGTTAAAGGAAAATGCTGC 1753
Db   1954 ACAGAATACCCCTTACAGCAAGAGGAGCTTAATTGACACCGTTTTAGTCAAGGAAAAGCTGC 2013

Qy   1754 GGCCAACACATCTTCAAAAAGAAGATGCTTAAAAAGAAATTGACTTCTACAGTTGTAATGAACTTATT 1813
Db   2014 AGCCAACACTCTTCAAAAACTCTCTGAGGAAAATTGACTTCCACGTTATATGAAAAACTTATT 2073

Qy   1814 TGTGGATAAGAGATATGAAGTATATATCCAACAGAGAGATGTTCAGGTGTCGTCACTGGAAGA 1873
Db   2074 TGTGGAAAAGAGAATATGAAGTTATTCCAACAGAAGACGTTTCAGGCTTGCATTGGGAAGA 2133

Qy   1874 ACAATTGAGGAGGTTGCAAGAAGAACGGAAACTTGTAAAGTGTGTATGGAGAAAAGAGTTTC 1933
Db   2134 GCAGTTGCCGGAGATTACAAGAGAACCAAACTGCAAAGTGTATGGGACAGAGAGGTTTC 2193

Qy   1934 TGTTGTATTTATTCCTTGTGGTCATCTGGTAGTATGCCAGGAATGTGCCCCTTCTCTAAG 1993

Query Match 51.9%; Score 1343.6; DB 3; Length 3151;
Best Local Similarity 76.2%; Pred. No. 3.5e-278;
Matches 1852; Conservative 0; Mismatches 499; Indels 79; Gaps 13;

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Qy    116  TTTGGTGGTAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAAGTGTTTTAGCTATCAA  175
Db    702  TGTGGTGGAGATCTAGTGTGTCCAAGTGGTGAGAAACTTCATCTGGAAGTTTAAGCGGGTCAG  761

Qy    176  ACAGCACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTC  235
Db    235  ...

Qy    236  GTATCAAAAACATTAAGAGTATAATTAATGGAAGCACGAATCTTGTCAGAGTTGGACAAACAG  295
Db    820  ACACCAAAAAACTTAAACGTATAAATGGAAGACGACAAATCTTGTCAAATTGGACAAAGGA  879

Qy    296  CAACAAAACAAAAATGAAGTAATGACTTTTCCTGTGAACTCAGAAGTCTCACATATTC  355
Db    880  GAGCGAAGAAAATGAAGTTTGACTTTTCCGTGTGAACTCTACCGAAGTCTACACATATTC  939

Qy    356  AACTTTCCCGCCCGGGGTGCCCTGTCTCAGAAAAGGAGTCTTGCTCGTCGTGCTGGTTTATTA  415
Db    880  ...

Qy    416  TACTGGTGTGAATGAACAAGTCAAAATGCTCTGTTGTTGGGCCTGATGCTGGATAACTGGAA  475
Db    1000  TACAGGTGTGAATGACAAAGTCAAGTGCTTCTGCTCGTGGGCCTGAATGTTGGATAACTGGAA  1059

Qy    476  ACTAGGAGACAGTGTCCCTATTCAAAAGCATAAAACAGCTATAATCCTTAGCGTTAGTCTTATTCA  535
Db    1060  ACAAGGGGACAGTCCTGTTGAAAAGCACACAGACAGTTCTATCCCAGCTCGCAGCTTTGTACA  1119

Qy    536  GAAATCTGGTTTCAGCTAGTCTGGGGATCCACCTCTAAGAATACGGTCTCCAAATGAGAAAACAG  595
Db    1120  GACTCTGCTTTCAGCCCAGTCTCGCAGTCTCCCATCTAAGGAATATATGTCTCCCTGTGAAAAAGTAG  1179

Qy    596  TTTTTGCACATTCATTATCTCCCCACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTC  655
Db    1180  ATTTGCACATTGGTCACCTTCTCAGGAACGAG----------GTGGGCATTCACTC  1221

Qy    656  CAGCCTTTCTCCAAAACCCTTCTTAATTCTAGAGCAGTTGAAGAGACATCTCTTCATCGAGGAC  715
Db    1222  CAACCTTGTGCTCTAGCCCTCTTAATTCTAGAGCAGTGGAAAGACTTC----TCATCAAGGAT  1278

Qy    716  TAAACCCCTACACGTTATTGCAATGAGTGACTGAAGAAGCCAGAGATTTCTTACCTACATATGTG  775
Db    1279  GGATCCCTTGCAGCTATGCCATGGACTGACTACACGAAGAGAGGCCAGAGATTTCTTACTTACAGTATGTG  1338

Qy    776  GCCATTAAACTTTTTTTGTCACCCATCAGAAATTGGCAAGAGAGCTCGGTTTTTATTATATAGGACC  835
Db    1339  GCCTTTAAGTTTTCTGTCACCAGCAGAGCTCGGCCCAGAGCCTGGTTCTTATTACATAGGGGCC  1398

Qy    836  TGGAGATTAGGGGGTTAGCCCTGCTTTGCCGTGTGGTGGGAGGCTCAGTAACTGGGGAAAGAACCCAAAGGA  895
Db    1399  TGGGAGCAGGGGTGGGGCCGTGTTTTCCCTGTGGGGAGAAACTGAGCAACTGGGAAGACCAAAGGA  1458

Qy    896  TGATGGCTATGTCAGAACACCGGAGGGCATTTTCCAACTGTCCATTTTTTGGGAAAAATTCTCT  955
Db    1459  TGATGCTAATGTCAGAGACGACCGCAGACGCATTTGTCCCACCTGTCAGCAGGGGGAAAAATACTTC  1518

Qy    956  AGAAACTTCTGAGGGGTTTTAGCATTTCAAAAATCTGAGCATGCAGCAACACATGCCAGCTCGAATGAG  1015
Db    1519  AGAAACACAGAGAGGTTTTAGTATGTATAACAAAATCTAAGTATGCACAGGCACACTCCTGCCTGAATGAG  1578

Qy    1016  AACATTTATGTACTGGGCCATTCTAGTGTTCCAGTTGTTCAGCGTGGCAGCTTGCAAGTGCTGG  1075
Db    1579  GACATTTCTGTTGACTGGGCCCACCTAGTGTTCCTGTTTTCAGCCCGAGCAGCTTGCAAGTGGCTGG  1638

Qy    1076  TTTTTATTTATGTGGGGGTGGCAAATGATGAGTCAAAATGCTTTTGTTGGTGATGGTGGCTTGAG  1135
Db    1639  ATTCTATTACCTGGATCGCAAATGATGAGTGCTTTTTGTTTGTTGTGATGGTGGCTTGAG  1698
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Qy   2054  AAAAATAGTCTATATTTTAACCTGCAATAAAAACGGTCTTTTAAAAATATTGTTGAACACTTTGA  2113
Db   2312  ------------GTGAAGAATGGTCTGTAAAGTATGTTGTTGGGACATCAGA  2347

Qy   2114  AGCCATCTAAAGTAAAAAGGGGAATTATGAGTTTTTCAATTAGTAACATTCAYTGTTCTAGT  2173
Db   2348  AGCTGTCAGAAACAAAGAAATGAAACTACTGA---TTTCAGCTCTCTCAGCAGGACACATTCTACT  2404

Qy   2174  CTGCTTTGGTACTAATAATCTTGTTTCTGAAAAAGATAGGTGTTCATATATATTTTAATCTTAAATC  2233
Db   2405  CTCTTTCAAGATTAGTTAATCTTGCTTTTATTGAAGGGTAGCTTGTATATTTTAAGCTTAGTC  2464

Qy   2234  TGTTTATTTACAAGGGGAAGATTAATGTTTGGTGAACTATATAATTAGTATGTAGTGTACCTA  2293
Db   2465  TG------TTGCAAGGGAAGGTCTATG-CTGTTGAGCTACANGGACTGGTTGTGTCGTTCCAGAG  2518

Qy   2294  AGGGAGTAGTGTCACTGCTTGTTATGCATCATTTCAGGANTTACTGGGATTTGTTGGTGTTCTT  2353
Db   2519  CAGGAGTTGGGATGCTTGCTGCTGTATGTC-----CTTCAGGA.TTCTTGGAT---TTGGAATT  2571

Qy   2354  TCAGAAAGCTTTGAATAACTAAAATTATAGTGGAAAAACAGAGAACTCTGG  2413
Db   2572  TGTGAAAGCTTTGGATTCAGGTGAGTGGGAGCTCAGAAA'CCTGAAAACCAGTGGGCTCTGG  2631

Qy   2414  AGTTCATCAGAGTTATGGGTGCCGAATTGTTCTTTGGGTGCTNTTCACTTGTGTGTTTTAAAATA  2473
Db   2632  TACTCA--GTAGTTAGGGTACCCTGTGCTTCTTGGGTGGCTNTTCCTTTCTG---GAAAATA  2686

Qy   2474  AGGATTTTTTCTTCTTTATTTCTCCCCCTAGTTTGTTGGTGAGAAA'ATCTCAATAAAAGTGCTTTAA  2533
Db   2687  AGGATTTTTCTGCTACTGGGTAAATATTTTCTGTTTGTGA;AAATATATAATTTAAAGTGTTTCT  2746

Qy   2534  AAAAAA  2539
Db   2747  TTTAAA  2752
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Qy   1994  AAAATGCCCTATTTGCAGGGGTATAATCAAGCAAAGTACTGTTCGTACATTTCTCTCTTAAAG  2053
Db   2254  GAAGTGCCCCATCTGCAGGGGGACAATCAAGCGGACCTGTH3CGCACATTTCTCATGA--  2311

Qy   2194  TATTGTGTTCATTCCGTGTGTGGTCATCTTAGTAGTAGTCTGCCAKGAATGTGCCCCTTCTCTAAG  2253
Db   ...
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Qy   1136  GTGTGGGGAAATCTGGACGTTAAGATGATCCATTGGGTAGTAGAACATGCCAAGGTGGTTTCCAAGGTGTGA 1195
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1699  ATGTTGGGGAAACCTGGAGATGAACCCCTGGATAGAACACGCCAAAATGGTTTCCAAGGTGTGA 1758

Qy   1196  GTTCTTGATACGGAATGAAAAGGCCAAGAGAGTTTGGTTGATTCAAGGTTAGATATCCTCA 1255
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1759  GTTCTTGATACGGATGAAGGGTCAGGAGTTTGATGAGAATATCCTCA 1818

Qy   1256  TCTTCTTCTTGAACAGCTGTTGTCAGATACCACTGGAGATAGAAAATGCTGACCC--- 1312
           |||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1819  TCTTCTTGAGCAGCTGTTGTCCACTTCAGACACCCCAGGAGAGAAAATGCTGACCCTAC 1878

Qy   1313  ---ACCAATTATTCATTTTGGACCTGGGAGAAAGTTCTTCAGAAGATGCTGTCATGGATGAA 1369
              ||||||||||||||||||||||||||||||||||||||||||||||
Db   1879  AGAGACAGTGGGTGCATTTTTGGCCCTGGAGAAAG---TTCGAAAGATGGTCGTCATGATGAG 1935

Qy   1370  TACACCGTGTGGTTAAAATCTGCCTTGGAAATGGGCTTTAATACGAGACCTGGGAAACAAAC 1429
           |||||||||||||||||||||||||||||||||||||||||||||||
Db   1936  CACGCCCTGTGGGTTAAAGCAGCCCTTGGAAATGGGCTTCAGCTAGGAGAGCCTGGGAGAGACAGAC 1995

Qy   1430  AGTTCAAAGTAAAATCCTGACAACTGGGAGGAGAGAACTATAAAACAGTTAATGATGATTATTGTGTC 1489
           |||||||||||||||||||||||||||||||||||||||||||||||
Db   1996  GGTTCAGCGCGGCAGAGATCCTGGCCACTGGTGTGAGAACTACACGGACCGTCAATGATATTGTCTC 2055

Qy   1490  AGCACTTCTAAATGCTGAGACGATGAAAAAAGAGGAGGAGGAGGAGGAAGCTGAAGA 1549
           ||||||||||||||||||||||||||||||||||||||||||||
Db   2056  AGTACATTTTGAAATGCTGAGACATGAAGAGAAGAAGAGGAGGAGGAAAGACAGACTGAAGA 2115

Qy   1550  AATGGCATCACGATGAGATTTGTCATTTAATTCGGAAGAACAGGAAATGGCTCTCTCTTCAACAATT 1609
           |||||||||||||||||||||||||||||||||||||||||||||||
Db   2116  GATGGCATCAGGTGAACTTATCACTGATTCGGAGAATAGAATGGCCCCTCTTCAACAGTT 2175

Qy   1610  GACATGTGTGCCTTCCTATCCTGGATAAATCTTCTTTAAAGGCCAATGTAATTAATAAAACAGGA 1669
           |||||||||||||||||||||||||||||||||||||||||||
Db   2176  GACACATGTCCTTCCTATCCTGGATAAATCTTCTTGAGGCCAGTGTAATTACAAAACAGGA 2235

Qy   1670  ACATGATTATTATTAAACAGAAAAAAACCAGATACCTTTACACAGGCGGAGAGAACTGATTGATAC 1729
           |||||||||||||||||||||||||||||||||||||||||||
Db   2236  ACATGATATTATTAGACACGAAAAACACAGATACCCTTACACAGCCAAGAGAGCTTAATTGACAC 2295

Qy   1730  CATTTTGGTTAAAGGAAAATGCTGCCGGCCCAACATCTTCAAAAACTGTCTAAAAGAAATTGA 1789
           |||||||||||||||||||||||||||||||||||||||||||
Db   2296  CGTTTTAGTTGCAAGGGAAAATGCTGCAGCCCACACATTCTCAAAAACTCTCTGAAAGAAAATTGA 2355

Qy   1790  CTCTACATTGTATAAGGACTTAATTATTTGGTGGATTAAGAATATGAAGTATATATTCCAACAGAAGA 1849
           |||||||||||||||||||||||||||||||||||||||||||
Db   2356  CTCCACGTTATATGTAGAAAACTTATTTGTGTAGGGAAAAGAATAATTGAAGTATATTCCAACAGAAGA 2415

Qy   1850  TGTTTCAGGTCTGTCACTGGAAGACAATTGAGGAGGTTGCAAGAGAAGACGAACTTGTAA 1909
           |||||||||||||||||||||||||||||||||||||||||||
Db   2416  CGTTTCAGGCTTGTCATTGGAAGAGCAGTTGCGGGAGATTACAAGAAGAACGAACTTGCAA 2475

Qy   1910  AGTGTGTATGGACAAAGAAGTTCTGTTGTGTATTTAATTCCTTGTGGGTCATCGGTAGTATG 1969
           |||||||||||||||||||||||||||||||||||||||||||
Db   2476  AGTGTGTATGGACAGAGAAGTTTTCATATTGTTTCCGTGTGGTGTTATCCTGAAGGAGTCTG 2535

Qy   1970  CCAGGAATGTGCCCCTTCTCCTAAGAGAAAATGCCCTATTTG(CAGGGGTGTATAATCAAGGGTAC 2029
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Db   2536  CCAGGAATGTTGCCCCCTTCTCCTAAAGGAAAGTGCCCCATCTGTCCAGGGGGACAAATCAAGGGGAC 2595

Qy   2030  TGTTCGTACATTTCTCTCTTAAAGAAAAATAGTCTATATATTTTAACCTGCATAAAAAGGTC 2089
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Db   2596  TGTGCGCGCACATTCTCTCATGA---------------GTGAAGAAATGGT 2629

Qy   2090  TTTAAAATATTGTTGAACACTTGAAGCCATCTAAAAGTAAAAAGGGAATTATGAGTTTTTTC 2149
           |||||||||||||||||||||||||||||||||||||||||||
Db   2630  CTGAAAAGTATTGTTGGACATCAGAGAAGCTGTCAGAACAAAGAAATGAACTACTGA---TTTC 2686

Qy   2150  AATTAGTAACATTCATGTTCTAGTCTGGCTTTGGTACTACTAATAATCTGTTTCTGAAAAGAT 2209
           |||||||||||||||||||||||||||||||||||||||||||
Db   2687  AGCTCTTCAGCAGACAAGTTACTCACTCTCTTCAAGATTAGTAATCTTGCTTATGAAAGGGT 2746

Qy   2210  GGTATCATATATTTAATCTTGTTATTACAAGGGAAGATTTATGTTGGGTGAAC 2269
           |||||||||||||||||||||||||||||||||||||||||||
Db   2210  GGTATCATATATTTAATCTTGTTATTACAAGGGAAGATTTATGTTGGGTGAAC 2269

Db   2747  AGCATTGTATATATTTAATTAAGCTTAGTTCTGTT------GCAAGGGGAAGGTCATATG-CTGTTGAGC 2800

Qy   2270  TATATTAGTTAGTGTATGTGTATGTACCTAAAGGGAGAGTAGTGTCACTCGTTGTTTATGCATCATTTCA 2329
           |||||||||||||||||||||||||||||||||||||||||||
Db   2801  TACAGGACCTGTGTCTGTTCCAGAGCAGGAGTGTTGGAATGCTTGGCGTGATGTC-----CTTCA 2856

Qy   2330  GGAGTTACTGGATTGTTGTTCTTTCAGAGAAAGCTTTGAAATACTAAATTATAGTGTAGAAA 2389
           |||||||||||||||||||||||||||||||||||||||||||
Db   2857  GGACTTCTTGGAT----TTGGAATTTGTGAAAAGCTTTGGATTCAGGTGGATCAGTGGAGCTCAG 2913

Qy   2390  AGAACTGGGAAACCAGGAACTCTCGAGTTCATCAGAGTTATGTGCCGAAATTGTCTTTTGGT 2449
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Db   2914  AAATCCTGAAAACCAGTGGCTCTGGTACTCA--GTAGTTAGGGTACCCTGTGCTTCTTGGT 2971

Qy   2450  GCTTTTCACTTGTTGTTTTAAAATAAGGATTTTTCTCTTATTTCTCCCCTAGTTTGTTGTGAG 2509
           |||||||||||||||||||||||||||||||||||||||||||
Db   2972  GCTTTTCCTTTCTG---GAAAATAAGGAGATTTTCTGCTACTGGTAAATATTTTCTGTTTG 3028

Qy   2510  AAACACTCTCAATAAAGTGCTTTAAAAAAAAA 2539
           |||||||||||||||||||||||||
Db   3029  TGAGAAATATATTAAAAGTGTTTCTTTTAAA 3058
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RESULT 13
US-08-800-929A-13
; Sequence 13, Application US/08800929A
; Patent No. 6133437
GENERAL INFORMATION:
 APPLICANT: Korneluk, Robert G
 APPLICANT: MacKenzie, Alexander E
 APPLICANT: Liston, Peter
 APPLICANT: Baird, Stephen
 APPLICANT: Tsang, Benjamin K
 APPLICANT: Pratt, Christine
 TITLE OF INVENTION: DETECTION AND MODULATION OF
 TITLE OF INVENTION: IAPS AND NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
 TITLE OF INVENTION: DISEASE
 NUMBER OF SEQUENCES: 17
 CORRESPONDENCE ADDRESS:
  ADDRESSEE: Clark & Elbing LLP
  STREET: 176 Federal Street
  CITY: Boston
  STATE: MA
  COUNTRY: USA
  ZIP: 02110
 COMPUTER READABLE FORM:
  MEDIUM TYPE: Diskette
  COMPUTER: IBM Compatible
  OPERATING SYSTEM: DOS
  SOFTWARE: FastSEQ for Windows Version 2.0
 CURRENT APPLICATION DATA:
  APPLICATION NUMBER: US/08/800,929A
  FILING DATE: 13-FEB-1997
  CLASSIFICATION: 424
 PRIOR APPLICATION DATA:
  APPLICATION NUMBER: 60/030,590
  FILING DATE: 14-NOV-1996
  APPLICATION NUMBER: 60/017,354
  FILING DATE: 26-APR-1996
 ATTORNEY/AGENT INFORMATION:
  NAME: Bieker-Brady, Kristina
  REGISTRATION NUMBER:
  REFERENCE/DOCKET NUMBER: 07891/009001
 TELECOMMUNICATION INFORMATION:
  TELEPHONE: 617-428-0200
  TELEFAX: 617-428-7045
  TELEX:
INFORMATION FOR SEQ ID NO: 13:
 SEQUENCE CHARACTERISTICS:
  LENGTH: 3151 base pairs
  TYPE: nucleic acid
  STRANDEDNESS: single

; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-800-929A-13

Query Match 51.9%;
Best Local Similarity 76.2%;
Matches 1852; Conservative 0; Mismatches 499; Indels 79; Gaps 13;

Score 1343.6; DB 3; Length 3151;
Pred. No. 3.5e-278;

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Qy    116 TTTGGTGGTAAAAATCTTAGTTGCATGTGAAGAAATTTCAIGTGAAGATGTTTAGCTATCAA 175
          |  |||| || |      |   |||||||  ||  ||||  | |  | ||| ||| |||||
Db    702 TGTGGTGGAGATCTAGTTGTCATGGAAAACTTCAICTGGAAGTTTAAGCGGTCAG 761

Qy    176 ACAGCACTGTCACCTACTCATGCACAAAAACTGCCTCCCCAAAGACTTTTCCCAGGTCCCTC 235
          |||||| | |||| || || |  |||||||| |||||| ||||| || ||| ||||||
Db    762 AAATACTATT--ACTACTCAIGGACAAAAAACTGTCTCCCAGAGACTCGGCGCCAAGGTACCTT 819

Qy    236 GTATCAAAACATTAAGAGTATAATAGGACACGAITCTTGTCAGATTGGACAAACAG 295
          ||| ||||| | ||||||| || |  ||  |||  |||| || ||||| |||||
Db    820 ACACCAAAAAACTTAAACGTATAAATGGAGAAGCACAAATCTTGTCAAAATTGGACAAAGGA 879

Qy    296 CAACAAACAAAAAAATGAAGTAATAGACTTTTCCCTGTGAACTCTACAGAAATGTCTACAITATTC 355
          ||  ||||  |||||| ||| |||| |||||||  |||| || | |  |||||||| |||||
Db    880 GAGCGAAGAAAAAAATGAAGTTTGACTTTGTGTGTGAACTCIACCGAAGTCTACAITATTC 939

Qy    356 AACTTTCCCGCCGCGGGGTGCCTGTCTCAGAAAGGAGTCTTJCTCGTGCGTGGTTTTTTTATTA 415
          ||||||||  | || ||||| |||||| || || ||||  |  |||| || || ||||| |||
Db    940 AGCTTTTTCCAGGGGAGTTCCGTTCTCGATAGCTTTCTGCTCTGCATGGCTTGGATAAAGTTC 999

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Qy    476 ACTAGAGGAGCAGTCCTATTCAAAAGCATAAACAGCTATATTCTTAGCTGTGTAGTTTATTCA 535
          ||||  |||||||||  |||||  |||| ||| |||||||| ||| || |  ||| |||||
Db   1060 ACAAGGGGACAGTCCTGTTGAAAAAGCACAGCAGTTCTTCTATCCCAGCTGCGACTTTGTACA 1119

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          ||||||||||||||| |||| | |||||||| |||||||  ||||| || |||||||
Db   1120 GACTCTGCTTTCAGCCAGTCTGCAGTCCAICTAAGAATAITGTCTCCTGTGAAAAGTAG 1179

Qy    596 TTTTGCACATTCATTATCTCCCACCTTGGGACATAGTAGCCITGTTCAGTGGTTCTTACTC 655
          |||| |||| || |  ||||||  ||| |   || |||||  |||||||||| || ||  |
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Qy    656 CAGCCTTTCTCCAAACCCTTCTTAATTCTAGAGACGCAGTGAACAICTCTTCATCGAGGAC 715
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Db   1222 CAACCTGTGCTTCTAGCCCTCTTAATTCTAGAGCAGTGGAAAGGACTTC---TCATCAAGGAT 1278

Qy    716 TAACCCCTACAGTTGTCAATGAGTAGTACTGAAGAAGCAGCAGAITTTCTTACCTACCATAGTG 775
          |||||| |||| |||| |||| || | |||| |  |||||| | |||||||| ||| || |||
Db   1279 GGATCCCCTGCAGCTAGCCATGAGGTACAGAGAGGCCACGAGATTTCTACTTTACTTACAGTATGTG 1338

Qy    776 GCCATTAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTCGTTTTTTAATATATAGGACC 835
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Db   1339 GCCTTTAAGTTTTCTGTCACCAGCAGAGGCTGGCTTCTCAITTCTATTACATAGGGGCC 1398

Qy    836 TGGAGATAGGGGTAGCCCTGCTTTGCCTGTGGGGAGCTCAGTGAACACAAAGGA 895
          ||| |||||||||  |||||| |||||| ||||  |||| ||||| ||||| ||
Db   1399 TGGGAACAGGGGTGGGGAAACTGACGAACTGGGAACCAAGGA 1458

Qy    896 TGATGCTAGTCAGAAACACCGGAGGGCAITTTCCCAACTGTCCAITTTTGGAAAAATTCTCT 955
          || |||| || ||||||| || | |||| |||||| ||||||  ||||| |||||||| ||
Db   1459 TGATGCTAITGTCAGAGCACCGCAGACAITTTCCCATTTCTGGAAAATACTTC 1518

Qy    956 AGAAAACTCTGAGGGTTTAGCATTCAAAATCTTAAGTATCCAGACACACTCTGCTCGATTGAG 1015
          |||| ||| |||||||||||| |||| |||| ||||| |  |||||| ||| | |||||||||
Db   1519 AGAAACACAGAGGGTTTAGTATATCAAAATCTAAGTATGGAGACACACTCTGCTGATTGAG 1578

Qy   1016 AACATTTAITGTACTGGCCATCTAGTGTTCAGTTCAGCCTTGCAAGTGCTGG 1075
          |  ||  || |||| ||  ||||| ||||||  | ||||||||  |||||||
Db   1579 GACATTTCTGTACTGGGCCACCTAGTGTTCCTGTTCAGCCTGAAGCTGTCTGG 1638

Qy   1076 TTTTTATTATGTTGGGTCGCAAATGCTTTTGTTGTTGTGAITGGTTGGCTTGAG 1135
          || || | | |  ||| |  ||||||  |||| |||||| || || |||| |||
Db   1639 ATTCTAITTACGITGGGAITCATCGCCAAGTGATGAITGCTTTTTGTTGTGATGGTGGCCTGAG 1698
```

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Db   1639 ATTCTAITTACGITGGGAITCATCGCCAAGTGATGAITGCTTTTTGTTGTGATGGTGGCCTGAG 1698

Qy   1136 GTGTTGGGAAATCTGGAGAITGATCCAITGGGTTAGAACATGCCAAGGTGTTGA 1195
          ||||||||||| | || ||| |||| ||||| || |||| |||||| |||| |
Db   1699 ATGTTGGGAACCTGGAGAITGACCCCTGGAITGAACACGCCAAATGGTTTCCAAGGTGTGA 1758

Qy   1196 GTTCTTGAITACGAITGAAAGGCCAAGAGTTGTTGATGGAITGAITGAGAITCAAGTAGAITATCCTCA 1255
          |||||||||  |||| ||  |||||||||||||| ||||| || ||||| |  |||| |||||||
Db   1759 GTTCTTGAITACGGATGAGGGGTCAGGGGTTGAGGAGTTTGAITGATGAITGACCC--- 1818

Qy   1256 TCTTCTTGAACAGCTGITGTCAAITCTTCAGAITTACCACTGGGAGAITGAGAAAAAITGCTGACCC--- 1312
          ||||||||||| |||  | ||||||| |||| ||  ||||| ||||| ||| || | ||| |||
Db   1819 TCTTCTTGAGCACGTGITTCCACTTCAGACACCCCAGGAGGAITGAAAATGCTGACCTAC 1878

Qy   1313 ---ACCAAITTATTCAITTTTGGACCTGGAGAAGGTTCTTCAGAITGAAGATGCTGCATGGAITGAA 1369
             |||| ||||||| |||||||||||||||  ||||| |||||| ||||  ||||||| |||||
Db   1879 AGAGACAGAGTGGITGCAITTTGGGCCCTGGGAAAG---TTCGAAAGATGITGCGTCGITCAITGGAITGAG 1935

Qy   1370 TACACCTGGITGGTTAAATCTGCCITTGGGAAAATGGCTTTAAITAGAITAGAITCCTGGAAAACAAAC 1429
          ||| |  |||| |||| ||||  ||| || |||||  |||| ||| |||||| |||| |||| |||
Db   1936 CACGCCTGITTGGGTTAAACGACCCTTGGGAAATGGGCTTCAGGTGGGTGAGACAGAC 1995

Qy   1430 AGTTCAAAGTAAAAITCCTGACAACTGGAGAITGAGAACTATAAAAAACAGITTAAITGAITAATTGTGTC 1489
          ||||||| | ||||||  | |||||||||||||||||||| | |||| ||||  || |  |||| | |
Db   1996 GGTTCAGCGGCGCAGAITCCGGCCACTGGTGAGAAGAACTTCAGGTAGGCGAAITGATAITTGTCTC 2055

Qy   1490 AGCACTTCTAAAITGCTGAAGAITGAAAAAAGAGAGGAGGAGGAGGAAAAAACAAGCTGAITGA 1549
          |||||||||||| || ||||  |||| | ||||| ||||||| || |  |||||||  |||||
Db   2056 AGTACTITTTGAITGCTGAAGAITGAGAGITGAGAGGAGGAGGAGGAAGAAGAAGAAGCAAGGA 2115

Qy   1550 AAITGGCAITCAGAITGAITGATTTTGTCATTAAITTCGGAAGGAACAGAITGGCTCTCTTTCAAITGA 1609
          || |||| | ||| ||| |||| || |||| || |||| |||||||||||| ||| |  || |||||
Db   2116 GAITGGCAITCAGGTGAITGACTTAITCACTGATTCGGAAGAITAGAAITGGCCCCTCTTGAITAITGA 2175

Qy   1610 GACAGTGTGCTTCCTAITCCCGAITAAIATCTTTTAAAGGCCAATGITAITTAAITTAAITAAAACAGGA 1669
          |||||||| |||||||| ||||| ||||||||| ||||||||  || |  ||||||||  ||||||||
Db   2176 GACACAGTGTCCITCCTAITCCCGAITAATCTTCTTCTTGAGGGCCAGTTAAITTACAGAAAACAGGA 2235

Qy   1670 ACAITGAITATTAITTATTAAAAITCAAAAAAITCAGAITAITCCTTTTACAAIGAITGAITGAITGAITAC 1729
          || ||| ||||| ||||||||| | |||||| ||||||   ||||  |||| |  ||| || |||||
Db   2236 ACAITGAITAITTAITTAGACAGAAAAACAITCAGAITACCCCTTACAAGGAGGCAITTAITTGACAC 2295

Qy   1730 CAITTTGGTTAAAGGAAAITGCTGCGGGCCCCAAACATCITCAAAAAITCGTGTCTAAAAGAAAAITTGA 1789
          |||||||||| ||| ||| ||| | | ||||| ||| ||| ||||| ||| ||  ||| |||||||
Db   2296 CGTTTTAGTCAAGGGAAAAGCTGCAITCTCCAGCCAACAITCTTCGAGAGGAAAAATTGA 2355

Qy   1790 CTCTACAITTGITAITAAGGAACTTAITTTGTGGAITAAGAITATATGGAGTATATTCCAACAGAAGA 1849
          || |||| | ||| || |||||||||||| ||||  || | |||| ||| | | ||||||||| ||
Db   2356 CTCCAGGTTAITATGAAAAACTTAITTGTGGAAAAGAITGAITATGGAGTATAITCCAAACAGAAGA 2415

Qy   1850 TGTTTCAGGTCTGTCACTGGGAAGAAGAAAACAAATTGAGGAGGAGGTTGCAAGAGAACGAACTTGTAA 1909
          || |||||| | |||||||| ||| || ||| |||||||  ||||||| ||| | |||| ||| |||
Db   2416 CGTTTCAGGCTCTGTCGTGITCAITGGAAGAGCAGITGGCGAGAITTACAAGAITGAACGAACTTGCAA 2475

Qy   1910 AGTGTGTATTGGAITCAIAAGAGAAGTTTCTGTTGTCAITTAITTTAITTCCCTGTTGGTCAITCTGGTAITG 1969
          |||||| |||||| || |||||||| || ||| | || ||||||||  ||| ||||| |||||||||
Db   2476 AGTGTGTATTGGAITCAITGGGAACAITGIAGAITCCGTGGTGITCAITCCTAITGTAITGTCTCAITGTATG 2535

Qy   1970 CCAGGAAITGTGGCCCCTTCTCTAITAAGAAAAIAITGCCCTAITTTGCAIGGGGGGGTATAITCAAGGGTAC 2029
          ||||||||||| ||||||||| || |||||| |||||  ||| ||| | | ||| ||| | |||||||
Db   2536 CCAGGAAAITGTGCCCCITCTGTTCTCTAAGAITAITGCCCAITCTGCAGGGGGGGACCAAITCAAGGGGAC 2595

Qy   2030 TGTTCGTACAITTTCTCTCTCTTAITAAAGAITAITTAGTCTAITATITTTTTAAACTGCAITAITAAAGGTC 2089
          ||||| |  |||||| || |||||||| || |  | | |   |||||   | ||| | |||||||||
Db   2596 TGTGCCGCACAITTTCTCCAITGA----------------------------GTGAITGAATGGT 2629

Qy   2090 TTTTAAAAITATTGGTGAAACAITCTTGAAGCCAITCTAAITGTAAAAGGAAITAAAGGGGAITTAITGAGITTTTTTC 2149
          ||| | | | || ||| |||||  |||||||  || | |||||  |||| |  |  | |  ||| ||
Db   2630 CTGAAAGTAITTGITGTGGAITCAITCAGAAGCGTGTCAGAAGAITGAACTACITGA---TTTC 2686

Qy   2150 AAITTAGTAAACAITTCAITGITTCTTCTAGTCTGCTITTGGTTTGITTACTGCITAAITAAITTACITTGTTTTTCTGAAAAAIGAT 2209
          ||||  |||||| ||||||   | ||| || ||| | |||| |  ||  | |  |   |  ||  ||| ||
Db   2150 —————————————————————————————————————————————— 2209
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Db   2687 AGCTCTTCAGCAGGACATTCTACTCTTCTTTCAAGATTAGTAATCTTGCTTTATGAAGGGT 2746

Qy   2210 GGTATCATATATTTAATCTGTTTATTACAAGGGAAGAGATTATTATTGTTTGGTGAAC 2269
Db   2747 AGCATTGTATATATTTAAGCTTAGTCTGTT-------GCAAGGGAAGGTCTATG-CTGTTGAGC 2800

Qy   2270 TATATATAGTTATGTGTATGTGTACCTAAGGGAGGTAGTAGTCACTGCTTGTTATGCATCATTTCA 2329
Db   2801 TACAGGACTGTGTCTGTTCCAGGACCAGGGATTGGGATGCTTGCTGTTGTTATGTC-----CTTCA 2856

Qy   2330 GGAGTTACTGGGATTTGTTGTTCTTTCAGAAACATTTTGAATTACTAAATTATATAGTGTAGAAA 2389
Db   2857 GGACTTCTTGGGAT---TTGGAATTTGTGAAAGCTTTGGATTCAGGTGAGGATGGGAAGCTCAG 2913

Qy   2390 AGAACTGGGAAAACCAGGGAACTCTCGGAGTTCATCAGAGTTATCTGGTGGTCCGAAATTGCTCTTTGGT 2449
Db   2914 AAAATCCTGAAAACCAGTGGCTCTCGGTACTCA--GTAGTTAGGGTAACCCTGTGCTCTTTGGT 2971

Qy   2450 GCTTTTCACTTGTGTTTTTAAAATAAAGGATTTTTCTCTTATTTCTCCCCCTAGTTTGTGAG 2509
Db   2972 GCTTTTCCTTTCTG---GAAAATAAGGGATTTTTTCTGGCTACTCGGTAAAATATTTTCTGTTTG 3028

Qy   2510 AAACATCTCAATAAAAGTGCTTTAAAAAAAAA 2539
Db   3029 TGAGAAATATATTAAAGTGTTTCTTTTTAA 3058

RESULT 14
US-09-617-053A-13
; Sequence 13, Application US/0961705A
; Patent No. 6300492
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: MacKenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAP's AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009003
; CURRENT APPLICATION NUMBER: US/09/617,053A
; CURRENT FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 08/800,929
; PRIOR FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 3151
; TYPE: DNA
; ORGANISM: Mus musculus
US-09-617-053A-13

Query Match  51.9%;  Score 1343.6;  DB 4;  Length 3151;
Best Local Similarity  76.2%;  Pred. No. 3.5e-278;
Matches 1852;  Conservative 0;  Mismatches 499;  Indels 79;  Gaps 13;

Qy    116 TTTGGGTGGTAAAAATCTTAGTTCATGTGAAGAAATTCATGTGAATGTTTTAGCTATCAA 175
Db   1759 GTTCCTTGATACGGATAGAAAGGCCAAGAGTTTGTTGTTGAATGAGAATTCAAGGTAGGATATCCTCA 1818

Qy    176 ACAGCACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTC 235
Db   1819 TCTTCTTTGAGCAGCTGTTGTTCCACTTCAGACACCCCAGGAGAGAAAAATGCTGACCCTAC 1878

Qy    236 GTATCAAAAACATTAAGAGTATAATGGAAGATAGCACGAGATTGGGACAAACAG 295
Db   1879 AGAGACAGTGGTGCATTTTGGCCCTGAGAAAG---TTGGAAAGATTGTCGTCATGATGAG 1935

Qy    296 CAACAAAACAAAAAATGAAGTATGACTTTTTCCTGTGAACTCTACGAATGTCTACATATTC 355
Db    880 GAGCGAAGAAAAATGAAGTTTGACTTTTTGTGTGTGAACTCTACGGAATGTCTACATATTC 939

Qy    702 TGTGGGTGGAGATCTAGTGTCCAAGTGGTGAGAAACTTCATCTGGAAGTTAAGCGGGTCAG 761
Db    762 AAATACTATT--ACTACTCATGCACAAAAACTGTCTCCCAGACTCGGGCCAAGGTACCTT 819

Qy   1136 GTGTTGGGAATTCGGAGTGAGATGACCCCTGAGAATGCACGCCAAATGGTTTCCAAGGTGTGA 1195
Db   1699 ATGTTGGGGACCTGGAGATGACCCCTGAGAATGCACGCCAAATGGTTTCCAAGGTGTGA 1758

Qy    776 GCCATTAACACTTTTTTGTCACCATCAGAATTGGCAAGAGAGTCTGGTTTTATTATATAGGACC 835
Db   1279 GGATCCCTGCAGCTATGCCATGCCATGAGTACAGAAGAGAGGCCAGATTTCTTACTTACAGTATGTG 1338

Qy   1016 AACATTTATGTTACTGCCCATCTAGTGTTGTTCCAGTTCAGCTGAGACGCTTGACAGTGCTGG 1075
Db   1519 AGAAACACAGAGGTTTAGTTATATCAAAATCTAAGTATGCAGAAGCACACTCTGCTCGATGAG 1578
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Qy       1430 AGTTCAAAGTAAAATCCTGACAACTGGAGAGAACTATAAAACAGTTAATGATATTGTGTC 1489
               |||||  |  | |||||| | ||||| |||||||| |  || || ||||||||||| ||
Db       1996 GGTTCAGCGGCAGATCCTGGCCACTGGTGAGAACTACAGGACCGTCAATGATATTGTCTC 2055

Qy       1490 AGCACTTCTAAATGCTGAAGATGAAAAAAGAGAGGAGGAGAAGGAAAAACAAGCTGAAGA 1549
              || |||| | |||||||||||||| | |||||| ||||||||||||| |||  |||||||
Db       2056 AGTACTTTTGAATGCTGAAGATGAGAGAAGAGAAGAGGAGAAGGAAAGACAGACTGAAGA 2115

Qy       1550 AATGGCATCAGATGATTTGTCATTAATTCGGAAGAACAGAATGGCTCTCTTTCAACAATT 1609
               |||||||||| ||| || ||| | ||||||||||| |||||||| ||||||||||| ||
Db       2116 GATGGCATCAGGTGACTTATCACTGATTCGGAAGAATAGAATGGCCCTCTTTCAACAGTT 2175

Qy       1610 GACATGTGTGCTTCCTATCCTGGATAATCTTTTAAAGGCCAATGTAATTAATAAACAGGA 1669
              ||||  ||| ||||||||||||||||||||| |  |||||| ||||||||  ||||||||
Db       2176 GACACATGTCCTTCCTATCCTGGATAATCTTCTTGAGGCCAGTGTAATTACAAAACAGGA 2235

Qy       1670 ACATGATATTATTAAACAAAAACACAGATACCTTTACAAGCGAGAGAACTGATTGATAC 1729
              |||||||||||||| ||| |||||||||||||| |||||||| ||||| || ||||| ||
Db       2236 ACATGATATTATTAGACAGAAAACACAGATACCCTTACAAGCAAGAGAGCTTATTGACAC 2295

Qy       1730 CATTTTGGTTAAAGGAAATGCTGCGGCCAACATCTTCAAAAACTGTCTAAAAGAAATTGA 1789
              | |||| || || ||||||||||| ||||||||||||||||||| ||| || ||||||||
Db       2296 CGTTTTAGTCAAGGGAAATGCTGCAGCCAACATCTTCAAAAACTCTCTGAAGGAAATTGA 2355

Qy       1790 CTCTACATTGTATAAGAACTTATTTGTGGATAAGAATATGAAGTATATTCCAACAGAAGA 1849
              ||| || || ||| | |||||||||||||| |||||||||||||||||||||||||||||
Db       2356 CTCCACGTTATATGAAAACTTATTTGTGGAAAAGAATATGAAGTATATTCCAACAGAAGA 2415

Qy       1850 TGTTTCAGGTCTGTCACTGGAAGAACAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAA 1909
               ||||||||  ||||| ||||||| || ||| |||| || ||||||||||||||||| ||
Db       2416 CGTTTCAGGCTTGTCATTGGAAGAGCAGTTGCGGAGATTACAAGAAGAACGAACTTGCAA 2475

Qy       1910 AGTGTGTATGGACAAAGAAGTTTCTGTTGTATTTATTCCTTGTGGTCATCTGGTAGTATG 1969
              |||||||||||||| ||| |||||| |||| || ||||| ||||||||||| ||||| ||
Db       2476 AGTGTGTATGGACAGAGAGGTTTCTATTGTGTTCATTCCGTGTGGTCATCTAGTAGTCTG 2535

Qy       1970 CCAGGAATGTGCCCCTTCTCTAAGAAAATGCCCTATTTGCAGGGGTATAATCAAGGGTAC 2029
              |||||||||||||||||||||||| || ||||| || |||||||| | ||||||||| ||
Db       2536 CCAGGAATGTGCCCCTTCTCTAAGGAAGTGCCCCATCTGCAGGGGGACAATCAAGGGGAC 2595

Qy       2030 TGTTCGTACATTTCTCTCTTAAAGAAAAATAGTCTATATTTTAACCTGCATAAAAAGGTC 2089
              ||| || ||||||||||| | |                             || || |
Db       2596 TGTGCGCACATTTCTCTCATGA--------------------------GTGAAGAATGGT 2629

Qy       2090 TTTAAAATATTGTTGAACACTTGAAGCCATCTAAAGTAAAAAGGGAATTATGAGTTTTTC 2149
               | ||| |||||||| |||   |||||  ||  ||  || || | | |  |||   ||||
Db       2630 CTGAAAGTATTGTTGGACATCAGAAGCTGTCAGAACAAAGAATGAACTACTGA---TTTC 2686

Qy       2150 AATTAGTAACATTCATGTTCTAGTCTGCTTTGGTACTAATAATCTTGTTTCTGAAAAGAT 2209
              |  |  ||      |  ||||| |||  ||    | || |||||||| ||    || | |
Db       2687 AGCTCTTCAGCAGGACATTCTACTCTCTTTCAAGATTAGTAATCTTGCTTTATGAAGGGT 2746

Qy       2210 GGTATCATATATTTAATCTTAATCTGTTTATTTACAAGGGAAGATTTATGTTTGGTGAAC 2269
               | ||  ||||||||| |||| ||||||      ||||||||| | ||||  || ||| |
Db       2747 AGCATTGTATATTTAAGCTTAGTCTGTT-----GCAAGGGAAGGTCTATG-CTGTTGAGC 2800

Qy       2270 TATATTAGTATGTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTATGCATCATTTCA 2329
              || |  | | ||| |||         ||||| | |   ||   |||    |     ||||
Db       2801 TACAGGACTGTGTCTGTTCCAGAGCAGGAGTTGGGATGCTTGCTGTATGTC----CTTCA 2856

Qy       2330 GGAGTTACTGGATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAATTATAGTGTAGAAA 2389
              ||| ||  |||||   |||   |||  |||||||||| || |         |      |
Db       2857 GGACTTCTTGGAT---TTGGAATTTGTGAAAGCTTTGGATTCAGGTGATGTGGAGCTCAG 2913

Qy       2390 AGAACTGGAAACCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCCGAATTGTCTTTGGT 2449
              | | |  ||||||||   ||||||   |||    ||||| ||| ||   |  |  |||||
Db       2914 AAATCCTGAAACCAGTGGCTCTGGTACTCA--GTAGTTAGGGTACCCTGTGCTTCTTGGT 2971

Qy       2450 GCTTTTCACTTGTGTTTTAAAATAAGGATTTTTCTCTTATTTCTCCCCCTAGTTTGTGAG 2509
              |||||||  || ||    |||||||||||||||||  || |  |     |  | |||  |
Db       2972 GCTTTTCCTTTCTG---GAAAATAAGGATTTTTCTGCTACTGGTAAATATTTTCTGTTTG 3028

Qy       2510 AAACATCTCAATAAAGTGCTTTAAAAAAAA 2539
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                |  | |  |  || ||    |||     |||
Db       3029 TGAGAAATATATTAAAGTGTTTCTTTTAAA 3058

RESULT 15
US-09-201-936-41
; Sequence 41, Application US/09201936
; Patent No. 6541457
; GENERAL INFORMATION:
;  APPLICANT: Korneluk, Robert G.
;  APPLICANT: MacKenzie, Alexander E.
;  APPLICANT: Baird, Stephen
;  APPLICANT: Liston, Peter
;  TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
;  TITLE OF INVENTION: PROBES, AND DETECTION METHODS
;  FILE REFERENCE: 07891/003003
;  CURRENT APPLICATION NUMBER: US/09/201,936
;  CURRENT FILING DATE: 1998-12-01
;  EARLIER APPLICATION NUMBER: 09/011,356
;  EARLIER FILING DATE: 1998-02-04
;  EARLIER APPLICATION NUMBER: PCT/IB96/01022
;  EARLIER FILING DATE: 1996-08-05
;  EARLIER APPLICATION NUMBER: 08/576,956
;  EARLIER FILING DATE: 1995-12-22
;  EARLIER APPLICATION NUMBER: 08/511,485
;  EARLIER FILING DATE: 1995-08-04
;  NUMBER OF SEQ ID NOS: 45
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 41
;   LENGTH: 2416
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-201-936-41

  Query Match             51.4%;  Score 1330.4;  DB 4;  Length 2416;
  Best Local Similarity   76.4%;  Pred. No. 2.2e-275;
  Matches 1816;  Conservative    0;  Mismatches  486;  Indels   74;  Gaps   12;

Qy        116 TTTGGTGGTAAAAATCTTAGTTCATGTGAAGAAATTTCATGTGAATGTTTTAGCTATCAA 175
              | ||||||  |     |       ||   ||||| ||||| || | |||| |||  |||
Db          2 TGTGGTGGAGATCTATTGTCCAAGTGGTGAGAAACTTCATCTGGAAGTTTAAGCGGTCAG 61

Qy        176 ACAGCACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTC 235
              | |      |  |||||||||| ||||||||| |||||| |||||   || |||| |||
Db         62 AAATACTATT--ACTACTCATGGACAAAACTGTCTCCCAGAGACTCGCCCAAGGTACCTT 119

Qy        236 GTATCAAAACA-TTAAGAGTATAATGGAAGATAGCACGATCTTGTCAGATTGGACAAACA 294
                | | ||| | ||||  ||||||||||  | ||||| ||||||||| ||||||||||
Db        120 ACACCCAAAAACTTAAACGTATAATGGAGAAGAGCACAATCTTGTCAAATTGGACAAAGG 179

Qy        295 GCAACAAACAAAAAATGAAGTATGACTTTTCCTGTGAACTCTACAGAATGTCTACATATT 354
                | | || |||||||||||| ||||||||| |||||||||||| |||||||||||||||
Db        180 AGAGCGAAGAAAAAATGAAGTTTGACTTTTCGTGTGAACTCTACCGAATGTCTACATATT 239

Qy        355 CAACTTTCCCCGCCGGGGTGCCTGTCTCAGAAAGGAGTCTTGCTCGTGCTGGTTTTTATT 414
              || |||| |||   || || ||||||||||| |||||||| ||||||||||| |||||||
Db        240 CAGCTTTTCCCAGGGGAGTTCCTGTCTCAGAGAGGAGTCTGGCTCGTGCTGGCTTTTATT 299

Qy        415 ATACTGGTGTGAATGACAAGGTCAAATGCTTCTGTTGTGGCCTGATGCTGGATAACTGGA 474
              |||| |||||||||||||| ||||| |||||||| |||||||||||| ||||||||||||
Db        300 ATACAGGTGTGAATGACAAAGTCAAGTGCTTCTGCTGTGGCCTGATGTTGGATAACTGGA 359

Qy        475 AACTAGGAGACAGTCCTATTCAAAAGCATAAACAGCTATATCCTAGCTGTAGCTTTATTC 534
              ||| ||| ||||||||| || ||||||| | |||| | ||||| ||||| |||||| | |
Db        360 AACAAGGGGACAGTCCTGTTGAAAAGCACAGACAGTTCTATCCCAGCTGCAGCTTTGTAC 419

Qy        535 AGAATCTGGTTTCAGCTAGTCTGGGATCCACCTCTAAGAATACGTCTCCAATGAGAAACA 594
              ||| |||| ||||||| ||||||   ||  | |||||||||| ||||||  ||| ||  |
Db        420 AGACTCTGCTTTCAGCCAGTCTGCAGTCTCCATCTAAGAATATGTCTCCTGTGAAAAGTA 479

Qy        595 GTTTTGCACATTCATTATCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACT 654
              | ||||||||||| | | |||        |                  ||||   | |||
Db        480 GATTTGCACATTCGTCACCTCTGGAACGAG------------------GTGGCATTCACT 521
```

Qy   655 CCAGCCTTCTCCCAAACCCTCTTAATTCTTAGAGCAGCAGTTGAAGACATCTCTTCATCGAGGA 714
           |||||||||||||||||||||||||||||||||||||||||||||||||| ||||||
Db   522 CCAACCTGTGCTTCTAGCCCTTCTTTAATTCTTAGAGCAGTTGGAAGACTTC---TCATCAAGGA 578

Qy   715 CTAACCCCTACAGTTATGCAAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCATATGT 774
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   579 TGGATCCCTGCAGCTATGCCATGAGTACAGAAGAGGCCAGATTTCTTACTTACAGTATGT 638

Qy   775 GGCCATTAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATAGAC 834
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   639 GGCCCTTTAAGTTTTTCGTCGTCACCAGCAGAGCCTTCTTATTACATAGGGGC 698

Qy   835 CTGGGAGATAGGGGTTAGCCTGCTTTGCCTTTGCCTGGGAAGCTCAGTAACTGGGAACCAAAGG 894
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   699 CTGGGAGACAGGGGTGGGCCTTGTTTTGCCTGTGGGTGGGTGGGGAAACTGAGCAACTGGGAACCAAAGG 758

Qy   895 ATGATGCTATGTCAGAACACCCGGAGGCACATTTTCCCAAACTGTCCATTTTTTTGGAAAAATTCTC 954
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   759 ATTATGCTATGTCAGAGCACCGCAGACATTTTCCCCACTGTCCATTTTCTTGGAAAATACTT 818

Qy   955 TAGAAACTCTGAGGTTTAGCATTTCAAATCTGAGCAATGCAGACACATCCAGCTCGAATGA 1014
           |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   819 CAGAAACACAGAGGTTTAGTATATCAAAATCTAAGTGATGCAGCACACTCCTGCTCGATTGA 878

Qy   1015 GAACATTTATGTACTGGCCATCTCAGTTGTTCCAGTTCAGCCCTGAGCGAGCAGCAGAAGTGCTG 1074
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   879 GGACACATTTCTGTACTGGCGCCACCTAGTGTTTCCTGTTCAGCCCGAGCAGCTTGCAAGTGCTG 938

Qy   1075 GTTTTTATTAATGTGGGGTCGGCAAATGAATGAGTCAAAATGCTTTTGTGTTTTGTTGATGGTGGCTTGA 1134
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   939 GATTCTATTACGTGGGATCGGCAATGATGATGTCAAGTCGCCTTGTTGTTGTGATGTGGGCTTGA 998

Qy   1135 GGTGGTTCGGGAAGTCTGGAGATGATCCATGGGCTAGAACATGCCAAGTGGTTTCCAAGGTGTG 1194
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   999 GATGTGGGGAACCTGGAGATGACCCCTGGAATAGAACAGCGCAAAATGGTTTTCCAAGGTGTG 1058

Qy   1195 AGTTCTTGATACGAAATGAAAAGGCCAAGAGAGTTTGTTGGTTGCATGAGATGTAGAGATATCCTC 1254
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Qy   1255 ATCTTCTTGAACACGTGTTGTCAACTTCACTCAGTGCGAGAAGAAAAAATGTGACCC-- 1312
           |||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1119 ATCTTCTTGAGCAGCTGTTGTCCACTTCAGACCAGTAGGAAGAAAAATGCTGACCCTA 1178

Qy   1313 -----ACCAATTATTCATTTTGGACCTGGAGAAAAGTTCTTCAGAAGATGCTGTCATGATGA 1368
                |||| |||||||||| ||||| ||||||||||||||||||||
Db   1179 CAGAGACAGTTGCATTTTGGCCCTGGAGGAAAG---TTCGAAAGATGTGTCATGATGATGA 1235

Qy   1369 ATACACCTGTGTTAAAACTGCCTTGGAAATGGGCTTTAAATAGAGACCTGGTGAAAACAAA 1428
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Db   1236 GCACGCCTGTGGGTTAAAGCAGCCTTGGAAATGGGCTTCAGTAGGAGCCTGGGTGACAGA 1295

Qy   1429 CAGTTCAAAGTAAAATCCTGACAACTGGAGAGAACTATAAAACAGTTAATGATAATTGTGT 1488
           |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1296 CGGTTCAGCGCGGCAGATCCTGGGCCACTGGGTGAGAGAACTACACGGACCTCAAATGATATATTGTCT 1355

Qy   1489 CAGCACTTCTAAATGCTGAAAGATGAAAAAGAGAGGAAAAACAAGCTGAAG 1548
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Db   1356 CAGTACTTTTGAATGCTGAAGATGAGAAAGAGGAAGAGGAAAACACAGACTGAAG 1415

Qy   1549 AAAATGGCATCAGAGATGATTTGTCATTAATTCGGAAGAACACAATAGGCTCTCTTTCAAACAAT 1608
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1416 AGATGGCATCAGGTGACTTATCACTCGATTAAATTCTTTTAAAGGCCAATGTAATTTAAAAACAGAGT 1475

Qy   1609 TGACATGTGCTTCCTTCCTAATCCTGGATAAATCTTCTTTAAAGGCCAGTGTTAAATTACAAAACAGG 1668
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Qy   1669 AACATGATAATTATTAAACAAAAAACACAGATACCCTTACCAAGCGAGAGAGACTGATTGATA 1728
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Qy   1909 AAGTGTGTATGGGACAAAAGAAGAGTTTCTCGTTTGTATTATTCCTTGTGGTTGGTGGTTGATTATT 1968
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Qy   2029 CTGTTCGGTACATTTCTCTCTTAAAGAAAAAATAGTCTATARTTTTAACCTGCATAAAAAGGT 2088
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Qy   2269 CTATATTAGTATAGTTATAGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTTATCGCATCATTTC 2328
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Db   2101 CTACAGGACGTGTCGTTCCAGAGCAGGAGATGCTTGCGTGTATGTC------CTTC 2156

Qy   2329 AGGAGTTACTCGGATTTGTTGTTCTTCAGAAAAGCTTTGAATACTAAATTATAGTGTAGAA 2388
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Qy   2389 AAGAACTGGAAAACCAGGAAACTCTGGGAGTTCATCAGAGTTATGGTGGCGGAATTGTCTTTTGG 2448
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GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: December 4, 2003, 14:22:05 ; Search time 815 seconds
(without alignments)
10558.035 Million cell updates/sec

Title: US-08-569-749-1
Perfect score: 2589
Sequence: 1 TCTAAGTAGTATCTTGGAAA.........AAAAAAAAAAAAAAAAAAA 2589

Scoring table: IDENTITY_NUC
                Gapop 10.0 , Gapext 1.0

Searched: 2201672 seqs, 166179599 residues

Total number of hits satisfying chosen parameters: 4403344

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB ID	Description
1	2589	100.0	2589	12	US-10-232-286-1	Sequence 1, Appli
2	2549.2	98.5	4614	12	US-09-814-353-19934	Sequence 19934, A
3	2535.8	97.9	3532	10	US-09-880-107-3354	Sequence 3354, Ap
4	2535.8	97.9	3732	10	US-09-974-592-7	Sequence 7, Appli
5	2531	97.8	2531	14	US-10-207-655-199	Sequence 199, App
6	2531	97.8	3496	14	US-10-153-668-337	Sequence 337, App
7	2514.8	97.1	2580	10	US-09-201-936-7	Sequence 7, Appli
8	1430.2	55.2	1435	8	US-08-464-588-1	Sequence 1, Appli
9	1430.2	55.2	1435	14	US-10-323-643-1	Sequence 1, Appli
10	1345.2	52.0	2862	12	US-10-232-286-13	Sequence 13, Appl
11	1343.6	51.9	3151	10	US-09-974-592-13	Sequence 13, Appl
12	1330.6	51.4	2416	10	US-09-201-936-41	Sequence 41, Appl
13	1157	44.7	3076	10	US-09-954-456-1635	Sequence 1635, Ap
14	1157	44.7	3076	10	US-09-954-531-16	Sequence 16, Appl
15	1157	44.7	3076	14	US-10-197-290-1	Sequence 1, Appl
16	1155.4	44.6	5857	12	US-10-247-671-18	Sequence 18, Appl
17	1155.2	44.6	3164	12	US-09-971-392-253	Sequence 253, App
18	1153.8	44.6	3165	12	US-10-141-618-5	Sequence 5, Appli
19	1153.8	44.6	2601	12	US-10-269-909-5	Sequence 5, Appli
20	1153.6	44.6	2601	12	US-10-232-286-3	Sequence 3, Appli
21	1152.2	44.5	6669	10	US-10-311-455-130	Sequence 130, App
22	1149	44.4	2676	10	US-09-201-936-5	Sequence 5, Appli
23	1078.4	41.7	2291	9	US-09-778-927A-21	Sequence 21, Appl
24	931.8	36.0	2676	12	US-09-201-936-39	Sequence 11, Appl
25	921.8	35.6	2450	10	US-09-201-936-39	Sequence 39, Appl
26	806.6	31.2	3734	12	US-10-353-461-7	Sequence 7, Appli
27	754.2	29.1	7049	12	US-10-311-455-129	Sequence 129, App
28	754.2	29.1	7049	12	US-10-240-452-6	Sequence 6, App
29	744	28.7	7049	12	US-10-311-455-127	Sequence 127, App
30	744	28.7	477	14	US-09-916-995-37203	Sequence 37203, A
31	475.4	18.4	477	14	US-10-102-524-591	Sequence 591, App
32	473.8	18.3	471	14	US-10-311-455-129	Sequence 129, App
33	406.4	15.7	460	10	US-09-960-352-8049	Sequence 8049, Ap
34	344.6	13.3	6681	12	US-10-240-452-3	Sequence 3, Appli
35	344.6	13.3	6681	12	US-09-960-352-8049	Sequence 128, App
36	305.2	11.8	460	10	US-10-311-455-128	Sequence 8049, Ap
37	296.4	11.4	6681	12	US-10-240-452-4	Sequence 128, App
38	296.4	11.4	6681	12	US-10-240-452-4	Sequence 4, Appli
39	274.2	10.6	419	10	US-10-102-524-4272	Sequence 4272, Ap
40	247.2	9.5	401	11	US-09-960-352-36214	Sequence 36214, A
41	243.4	9.4	565	12	US-10-029-386-125570	Sequence 125570,
42	195.8	7.6	255	14	US-10-102-524-906	Sequence 906, App
43	149.6	5.8	240	14	US-09-796-692-3493	Sequence 3493, Ap
44	149.6	5.8	240	14	US-10-040-862-3493	Sequence 3493, Ap
45	148	5.7	240	10	US-09-796-692-6687	Sequence 6687, Ap

ALIGNMENTS

RESULT 1
US-10-232-286-1
; Sequence 1, Application US/10232286
; Publication No. US20030143579A1
; GENERAL INFORMATION:
; APPLICANT: Rothe, Mike
; APPLICANT: Goeddel, David V
; TITLE OF INVENTION: INHIBITORS OF APOPTOSIS
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/232,286
; FILING DATE: 30-Aug-2002
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/569,749
; FILING DATE: <Unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: Brezner, David J.
; REGISTRATION NUMBER: 24,774
; REFERENCE/DOCKET NUMBER: A-62464/DJB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415)781-1989
; TELEFAX: (415)398-3249
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2589 base pairs
; TYPE: nucleic acid

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;
;
;                    STRANDEDNESS: single
;                    TOPOLOGY: linear
;                    MOLECULE TYPE: cDNA
US-10-232-286-1      SEQUENCE DESCRIPTION: SEQ ID NO: 1:

Query Match          100.0%;   Score 2589;   DB 12;   Length 2589;
Best Local Similarity 100.0%;  Pred. No. 0;
Matches 2589;   Conservative   0;   Mismatches   0;   Indels   0;   Gaps   0;
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Qy     1  TCTAAGTAGTATCTTGGGAAATTCAGAGAGATACTCATCCTACCTGAATATAAAACTGAGAT    60
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Qy    61  AAAATCCAGTAAAGAAGAGTGTAGTAGTAAAATTCTACATAAGAGTCTATCATTGATTCTTTTGG   120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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(Sequence alignment continues — paired Query (Qy) / Database (Db) nucleotide rows with 100% identity, positions 1–2040, matching bars shown between each pair.)

Qy   121 TGGTAAAAAATCTTAGTTCAITGTGAAGTGAAGAAAATTCATGTGAAGTGTTTTTTACTTATTCAAAACAGC   180
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Qy   181 ACTGTCACCTACTCAITGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTCGTATC   240
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Db   781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGGTTTTTATTATATAGGACCTGGAG   840

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RESULT 2
US-09-814-353-19934
Sequence 19934, Application US/09814353
Publication No. US20030165831A1
GENERAL INFORMATION:
APPLICANT: Lee, John
APPLICANT: Thompson, Pamela
APPLICANT: Lillie, James
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY OF OVARIAN CANCER
FILE REFERENCE: MRI-006B
CURRENT APPLICATION NUMBER: US/09/814,353
CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/191,031
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: US 60/207,124
PRIOR FILING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: US 60/211,940
PRIOR FILING DATE: 2000-06-15
PRIOR APPLICATION NUMBER: US 60/216,820
PRIOR FILING DATE: 2000-07-07
PRIOR APPLICATION NUMBER: US 60/220,661
PRIOR FILING DATE: 2000-07-25
PRIOR APPLICATION NUMBER: US 60/257,672
PRIOR FILING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 22037
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 19934
LENGTH: 4614
TYPE: DNA
ORGANISM: Homo sapiens

; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 4595, 4613, 4614
; OTHER INFORMATION: n = A,T,C or G
US-09-814-353-19934

Query Match   98.5%;   Score 2549.2;   DB 12;   Length 4614;
Best Local Similarity   99.6%;   Pred. No. 0;
Matches 2554;   Conservative   0;   Mismatches   9;   Indels   0;   Gaps   0;

Qy   1 TCTTAAGTAGTATCTTGGAAATTCAGAGAGATACTATCCTACCTGAATATAAAACTGAGAT   60
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Db   2036 TCTTAAGTAGTATCTTGGAAATTCAGAGAGATACTATCCTACCTGAATATAAAACTGAGAT   2095

Qy   61 AAATCCAGTAAAGAAAAGGTAGTAAAATTCTACATAAGAGTCTATCATTGATTTCTTTTGG   120
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Db   2096 AAATCCAGTAAAGAAAAGGTAGTAAAATTCTACATAAGAGTCTATCATTGATTTCTTTTTG   2155

Qy   121 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTCATGTGAAGTGTTTTAGCTATCATCAAAC   180
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Db   2156 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTCATGTGAAGTGTTTTAGCTATCAAACAGT   2215

Qy   181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAGACTTTTCCCAGTTCCCTCGTATC   240
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2216 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAGACTTTTCCCAGTTCCCTCGTATC   2275

Qy   241 AAAACATTAAGAGTATAATGAAGATAGCACAGATCTTGTCAGATTGGACAAAACAGCAACA   300
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Qy   301 AACAAAAAATGAAGTATGACTTTTCCTGTGAAACTCTACAGAAGTCTACATATTCAAACTT   360
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Db   2336 AACAAAAAATGAAGTATGACTTTTCCTGTGAAACTCTACAGAAGTCTACATATTCAAACTT   2395

Qy   361 TCCCCGCCGGGGTGCCTGTCTCAGAAAGGAGTCTTGCTCGTGCGTGGTTTTTATTATACTG   420
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Db   2396 TCCCCGCCGGGGTGCCTGTCTCAGAAAGGAGTCTTGCTCGTGCGTGGTTTTTATTATACTG   2455

Qy   421 GTGTGAATGACAAGGTCAAATGCTTCGTTGTTGTGGGCCTGAGTGCTGGATAACTGGAAAC   480
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Db   2456 GTGTGAATGACAAGGTCAAATGCTTCGTTGTTGTGGGCCTGAGTGCTGGATAACTGGAAAC   2515

Qy   481 GAGACAGTCCTATTCAAAAGCATAAAACAGCTATATCCTAGCGTAGCTTATTCAGAATC   540
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Db   2516 GAGACAGTCCTATTCAAAAGCATAAAACAGCTATATCCTAGCGTAGCTTATTCAGAATC   2575

Qy   541 TGGTTTCAGCTAGTCTGGGGATCCACCTCTAAGAATAGCTCTCCAAATGAGAAACAGTTTG   600
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Qy   601 CACATTCATTATCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC   660
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Db   2636 CACATTCATTATCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC   2695

Qy   661 TTTCTCCAAACCCTCTTAATTCTAGAGACAGTTGAAGACAATCTCTTCATCGAGGACTAACC   720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2696 TTTCTCCAAACCCTCTTAATTCTAGAGACAGTTGAAGACAATCTCTTCATCGAGGACTAACC   2755

Qy   721 CCTACAGTTATGCAATGAGTAGTACTGAAGAAGCCAGATATTCTTACCTACCATATGTGGCCAT   780
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2756 CCTACAGTTATGCAATGAGTAGTACTGAAGAAGCCAGATATTCTTACCTACCATATGTGGCCAT   2815

Qy   781 TAACTTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTATTATTATAGGAGACTTGGG   840
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2816 TAACTTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTATTATTATAGGAGACTTGGG   2875

Qy   841 ATAGGGTAGCCTCGCTTTGGGGAAGCTCAGTAACTGGGAAACCAGGATGATG   900
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Db   2876 ATAGGGTAGCCTCGCTTTGGGGAAGCTCAGTAACTGGGAAACCAGGATGATG   2935

Qy   901 CTATGTCAGAACACCGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAAATTCTCTAGAAA   960
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Db   2936 CTATGTCAGAACACCGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAAATTCTCTAGAAA   2995

Qy   2041 TTCTCTCTTAAAGAAAAATAGTCTATATTTAACCTGCA:AAAAAGGTCTTTAAAATATT   2100
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Qy   2101 GTTGAACACTTGAACCGCCATCTAAAAGTAAAAAGTAAAAAGGGAAATTA:GAGTTTTTCAATTAGTAGTAAACA   2160
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Db   2101 GTTGAACACTTGAACCGCCATCTAAAAGTAAAAAGTAAAAAGGGAAATTAAGAGTTTTTCAATTAGTAGTAAACA   2160

Qy   2161 TTCATGTTCTTAGTCTGCTTGGGTACTTAATAATCCTTGTTTCTGAAAAGATGGTATCATATA   2220
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Db   2161 TTCATGTTCTTAGTCTGCTTGGGTACTTAATAATCCTTGTTTCTGAAAAGATGGTATCATATA   2220

Qy   2221 TTTTAAATCTTAAATCTGTTTATTTATTACAAGGGAAGAGTTTATGTTGGTGAACTATATTAGTTAT   2280
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Db   2221 TTTTAAATCTTAAATCTGTTTATTTATTACAAGGGAAGAGTTTATGTTGGTGAACTATATTAGTTAT   2280

Qy   2281 GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTATGCATCAATTCAGGAGTTACTGG   2340
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Db   2281 GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTATGCATCAATTCAGGAGTTACTGG   2340

Qy   2341 ATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAAATTATATGTTGTAGAAAACTGGAAA   2400
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Db   2341 ATTTGTTGTTCTTTCAGAAAGCTTTGAATACTAAAATTATATGTTGTAGAAAACTGGAAA   2400

Qy   2401 CCAGGAAACTCTGGAGTTCATCAGAGTTATGGTGCCGGAATTGTTTCTTTGGTGTCTTTTTTCACTT   2460
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Db   2401 CCAGGAAACTCTGGAGTTCATCAGAGTTATGGTGCCGGAATTGTTTCTTTGGTGTCTTTTTTCACTT   2460

Qy   2461 GTGTTTTAAAATAAGGATTTTTTCTCTTATTTCTCCCCTAGTTTGGTGAGAAAACATCTCAA   2520
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Db   2461 GTGTTTTAAAATAAGGATTTTTTCTCTTATTTCTCCCCTAGTTTGGTGAGAAAACATCTCAA   2520

Qy   2521 TAAAGTGCTTTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAACTT   2580
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Db   2521 TAAAGTGCTTTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA   2580

Qy   2581 AAAAAAAAAA   2589
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Db   2581 AAAAAAAAAA   2589

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Qy    961  CTCTGAGGTTTAGCATTTCAAATCTGAGCATGCAGACA(ATGCAGCTCGAATGAGAACAT 1020
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Qy   1021  TTATGTACTGGGCCATCTACTAGTTGTTCCAGTTCAGCGACTTGCAAGTGCTCGGTTTTT 1080
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Db   3056  TTATGTACTGGGCCATCTACTAGTTGTTCCAGTTCAGCGACTTGCAAGTGCTCGGTTTTT 3115

Qy   1081  ATTATGTGGGTCGGCAATGATGATGATCAAATGCTTTTGTTGTGTAGGGTGGCCTTGAGG 1140
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Db   3116  ATTATGTGGGTCGGCAATGATGATGATCAAATGCTTTTGTTGTGTAGGGTGGCCTTGAGG 3175

Qy   1141  GGGAAATCTCGGAGATGATCCATGGGTAGGAACATGCCAAGTGGTTTCCAAGGTGTTCT 1200
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3176  GGGAAATCTCGGAGATGATCCATGGGTAGGAACATGCCAAGTGGTTTCCAAGGTGTTCT 3235

Qy   1201  TGATACGAATGAAAGGGCCAAGAGTTTGTTGATGGATTCAAGGTAGGATATCCTCATCTC 1260
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3236  TGATACGAATGAAAGGGCCAAGAGTTTGTTGATGGATTCAAGGTAGGATATCCTCATCTC 3295

Qy   1261  TTGAACAGCTGTTGTTCAACTTCAGATACCCACTGGGAGGAAGGAAAATGCTAGCCACCA 1320
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3296  TTGAACAGCTGTTGTTCAACTTCAGATACCCACTGGGAGGAAGGAAAATGCTAGCCACCA 3355

Qy   1321  TTCATTTTGGACCTGGAGAAAGTTCCTTCAGATGCTGT(CAGATGAATACACCTGTGG 1380
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Qy   1381  TTAAAATCTGCCTTGGGAAATGGGCCTTTAAATAGGAACCTGGGATGAAACAAGTTCAAAGTA 1440
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Db   3416  TTAAATCTGCCTTGGGAAATGGGCTTTAAATAGGAACCTGGGATGAAACAAGTTCAAAGTA 3475

Qy   1441  AAATCCTGACAACTGGAGAGAACTATAAAAACAGTTAATGX(TATTGTGTCAGCACTTCTAA 1500
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Qy   1501  ATGGTGAAGATGAAAAAAGAGAGGAAGAAAAACAAGCTGAGCAAGAAAATGCATCAG 1560
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Db   3536  ATGCTGAAGATGAAAAAAGAGAGGAAGAAAAACAAGCTGAGCAAGAAAATGCATCAG 3595

Qy   1561  ATGATTTGTCATTAATTCGGAAGAACAGATAGGCTCTCTTCTTTCAACAATTGACATGTGTGC 1620
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Qy   1741  AAGGAAAATGCTGCGGCCAACATCTTCAAAAAACTGTCTAAAAGAAAATTGACTCTACATTGT 1800
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Db   3776  AAGGAAAATGCTGCGGCCAACATCTTCAAAAAACTGTCTAAAAGAAAATTGACTCTACATTGT 3835

Qy   1801  ATAAGAACTTATTTGTGGATAAGAAATAATGAAGTATATTCCACACAGAAGAAGAGAGATGTTTCAGGTC 1860
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Db   3836  ATAAGAACTTATTTGTGGATAAGAAATATTCCACAGAAGAGATGTTTCAGGTC 3895

Qy   1861  TGTCACTCGAAGAACAAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTGTATGG 1920
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Db   3896  TGTCACTCGAAGAACAAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTGTATGG 3955

Qy   1921  ACAAAGAAGTTTCTGTTGTATTTAATTCCTTGTGTGGTCATCTCGGTTAGTAATGCCAGGAAATGTG 1980
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Db   3956  ACAAAGAAGTTTCTGTTGTATTTAATTCCTTGTGTGGTCATCTCGGTTAGTAATGCCAGGAAATGTG 4015

Qy   1981  CCCCTTCTCTTAAGAAAAATGCCTATTTGCAGGGGTACTGCTGTTCGTCGTACAT 2040
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4016  CCCCTTCTCTTAAGAAAATGCCTATTTGCAGGGGTACTGCTGTTCGTCGTACAT 4075

Qy   2041  TTCTCTCTTAAAAGAAAAATAGTCTATATATTTTAACCTGCATCAAAAAAGGTCTTTAAAATATT 2100
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Db   4076  TTCTCTCTTAAAAGAAAAATAGTCTATATATTTTAACCTGCATCAAAAAAGGTCTTTAAAATATT 4135

Qy   2101  GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGAATTATATGAGTTTTTTCAATTAGTAACA 2160
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Db   4136  GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGAATTATATGAGTTTTTTCAATTAGTAACA 4195

Qy   2161  TTCATGTTCTAGTCTGCTTTGGTACTAAATAAATCTTGTTTCTTGAAAAAGATGGTATCATTATA 2220
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Db   4196  TTCATGTTCTAGTCTGCTTTGGTACTAAATAAATCTTGTTTCTTGAAAAAGATGGTATCATTATA 4255

Qy   2221  TTTAAACTTAAACTGTTTATTTTACAAGGAGAAGAGATTTATGTTGTTGGTGAACTATATTAGTAT 2280
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Db   4256  TTTAAATCTTAAACTGTTTATTTTACAAGGAGAAGAGATTTATGTTGTTGGTGAACTATATTAGTAT 4315

Qy   2281  GTATGTGTACCTTAAGGGAGAGTAGTGTCACTCGCTTGTTATGCATCATTTCAGGAGTACTGG 2340
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Qy   2341  ATTTGTTGTTCTTTCAGAAAAAGCTTTTGAATACTAAAATTATAATTATAGTGTAGAAAAAGAACTGGAAA 2400
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Qy   2521  TAAAGTGCTTTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2563
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Db   4556  TAAAGTGCTTTAAAAAAAAAAAAAAAAAAAAACAAAAAAAAACACAAAAAAAA 4598
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RESULT 3
US-09-880-107-3354
; Sequence 3354, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
; APPLICANT: Horne, Darci T.
; APPLICANT: Vockley, Joseph G.
; APPLICANT: Scherf, Uwe
; APPLICANT: Gene Logic, Inc.
; TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
; FILE REFERENCE: 44921-5028-WO
; CURRENT APPLICATION NUMBER: US/09/880,107
; CURRENT FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/211,379
; PRIOR FILING DATE: 2000-06-14
; PRIOR APPLICATION NUMBER: US 60/237,054
; PRIOR FILING DATE: 2000-10-02
; NUMBER OF SEQ ID NOS: 3950
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 3354
; LENGTH: 3532
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: Genbank Accession No. US20020142981A1 U37547
US-09-880-107-3354

Query Match 97.9%; Score 2535.8; DB 10; Length 3532;
Best Local Similarity 99.7%;
Matches 2540; Conservative 0; Mismatches 7; Indels 0; Gaps 0;

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Qy    1  TCTAAGTAGTATCTTGAAATTCAGAGAGATACTCATCCTCTACCTGAATATAAACTGAGAT 60
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Qy    181 ACTGTCACCTTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTCGTATC 240
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Db   2466 ATGCTGAAGATGAAAAAGAAGAGGAGGAGGAAAAACAAGCTGAAGAAATGGCATCAG 2525

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          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy    541 TGGTTTCAGCTAGTCTGGGGATCCACCTCTCCAAATACGTCTCCAAATGAGAAAACAGTTTTG 600
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Qy   1621 TTCCTATCCTGGATAATCTTTTAAAAGGCCAAATGTAATTAATTAATAAACAGGAACCATGATTATTA 1680
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Db   2586 TTCCTATCCTGGATAATCTTTTAAAGGCCAAATGTAATTAATTAATAAACAGGAACCATGATTATTA 2645

Qy    601 CACATTCATTATCTCCCACCTTGGAACATAGTTAGCTTGTTTCAGTGGTTGTTCTTACTCCAGCC 660
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Qy   1681 TTAAAACAAAAAACACAGATAACCTTTACACAGCGAGAGCAACTGATTGATAACCATTTTTGGTTA 1740
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2646 TTAAAACAAAAAACACAGATAACCTTTACACAGCGAGAGCAACTGATTGATAACCATTTTTGGTTA 2705

Qy    661 TTTCTCCAAACCCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATGAGGACTAACC 720
          |||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1626 TTTCTCCAAACCCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATGAGGACTAACC 1685

Qy   1741 AAGGAAAATGCTGCGGGCCCAACACATCTTCAAAAAACTGTCTAAAAAAGAAAATTGACTCTTACATTGT 1800
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2706 AAGGAAAATGCTGCGGGCCCAACATCTTCAAAAAACTGTCTAAAAAAGAAAATTGACTCTTACATTGT 2765

Qy    721 CCTACAGTTATGCAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCATATGTGGCCAT 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1686 CCTACAGTTATGCAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCATATGTGGCCAT 1745

Qy   1801 ATAAGAACTTATTTTGTGGATAAGAATATGAAGTTACCAGATAGAAGAGAATTGCAGGGTC 1860
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2766 ATAAGAACTTATTTTGTGGATAAGAATATGAAGTTACCAGATAGAAGAGAATTGCAGGGTC 2825

Qy    781 TAACTTTTTTTGTCACCATCAGAATTGGCACCAAGAACACCGGAGGCATTTTCCCACTGTCCATTTT 840
          |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1746 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATAGGACCTGGAG 1805

Qy   1861 TGTCACTGGGAAGACAATTGAGGAGGGTTGCAAGAAGAAGGTTGCAAGAACTTGTTAAAGTGTGTATGG 1920
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2826 TGTCACTGGGAAGACAATTGAGGAGGGTTGCAAGAAGAAGGTTGCAAGAACTTGTTAAAGTGTGTATGG 2885

Qy    841 ATAGGGGTAGCCCTGCTTTGCCTCGTGGTGGGGAAGCTCAGTGAACTCGGAAGGGATGAGTG 900
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1806 ATAGGGGTAGCCCTGCTTTGCCTGTGGTGGGGAAGCTCAGTGAACTGGGAAGGGATGAGTG 1865

Qy   1921 ACAAAGAAGTTCTCTGTTGTTATTTATTTCCTTGTGATTCTCTGGTAGTTAATGCCAGGAATGTG 1980
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2886 ACAAAGAAGTTCTCTGTTGTTATTTATTTCCTTGTGATTCTCTGGTAGTATGCCAGGAATGTG 2945

Qy    901 CTATGTCAGAACACCGGAGGCATTTCCCACTGTCCATTTTTGGAAAAAATTCTCTAGAAA 960
          |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1866 CTATGTCAGAACACCGGAGGCATTTCCCACTGTCCATTTTTGGAAAAAATTCTCTAGAAA 1925

Qy   1981 CCCCTTCTCTAAGAAAAATGCCCTATTTGCAGGGGTATAATCAAAGGGTACGTTCGTTACAT 2040
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2946 CCCCTTCTCTAAGAAAAATGCCCTATTTGCAGGGGTATAATCAAAGGGTACTGTTCGTTACAT 3005

Qy    961 CTCTGAGGTTTAGCATTTCAAATCTGAGCATTGCAGACACATGCAAGCTCGAAATGAGAAACAT 1020
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1926 CTCTGAGGTTTAGCATTTCAAATCTGAGCATTGCAGACACATGCAAGCTCGAAATGAGAAACAT 1985

Qy   2041 TTCTCTCTTAAAGAAAAAATAGTCTATATATTTTAACCTGCATAAAAAGTTTCTTTAAAATATT 2100
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3006 TTCTCTCTTAAAGAAAAATAGTCTATATATTTTAACCTGCATAAAAAGTTTCTTTAAAATATT 3065

Qy   1866 CTATGTCAGAACACCGGAGGCATTTCCCAACTGTCCATTTTTGGAAAAAATTCTCTAGAAA 1925
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1866 CTATGTCAGAACACCGGAGGCATTTCCCAACTGTCCATTTTTGGAAAAAATTCTCTAGAAA 1925

Qy   2101 GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGGAATTATGAGTTTTTCAAGTTGTCATATA 2160
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3066 GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGGAATTATGAGTTTTTCAAGTTGTCATATA 3125

Qy   1021 TTATGTACTGGGTTCCAGTTCAGGCTGCCAAGTGCTGCTGGTTTTT 1080
          |||||||||||||||||||||||||||||||||||||||||||||
Db   1986 TTATGTACTGGGTTCCAGTTCAGGCTGCCAAGTGCTGCTGGTTTTT 2045

Qy   2161 TTCATGTTCTAGTCTCTGCTTTGGGTACTAATATAATCTTGTTTTCTGAAAAAGGATGTCATATA 2220
          ||||||||||||||||||||||||||||||||||||||||||||||||
Db   3126 TTCATGTTCTAGTCTCTGCTTTGGGTACTAATATAATCTTGTTTTCTGAAAAAGGATGTCATATA 3185

Qy   1081 ATTATTGTGGGGTCGCAAATGAGATGTCAAAATCGCTTTTGTTGTGTTTATGGTGCTTGAGGGTGTT 1140
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2046 ATTATTGTGGGGTCGCAAATGAGATGTCAAAATCGCTTTTGTTGTGTTTATGGTGCTTGAGGGTGTT 2105

Qy   1141 GGGAATCTGGAGATGATCCATGGGGTAGAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 1200
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1141 GGGAATCTGGAGATGATCCATGGGGTAGAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 1200

Qy   2221 TTTAATCTTAATCTGTTTTATTTTACAAAGGGAAAGATTTAGTTTGGGTGAACTATATTAGTTAT 2280
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3186 TTTAATCTTAATCTGTTTTATTTTACAAGGGAAAGATTTAGTTTGGGTGAACTATATTAGTTAT 3245
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Qy   2281 GTATGTGTACCTAAGGGAGTAGTGTGTCACTGCTTGTTTATTCATCATTCAGGAGTTACTGG 2340
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3246 GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTATGCATCATCATTCAGGAGTTACTGG 3305

Qy   2341 ATTTGTTGTTCTTTCAGAAGAGCTTTGAATACTAAATTA TAGTGTAGAAAGAGAACTGGAA 2400
          ||||||||||||||||||||||||||||||||||||||    |||||||||||||||||||
Db   3306 ATTTGTTGTTCTTTCAGAAAAGCTTTGAATACTAAATTATATAGTGTAGAAAGAGAACTGGAAA 3365

Qy   2401 CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGGTGGCCGAATTGTCTTGGTGCTTTTCACTT 2460
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3366 CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCCGAATTGTCTTGGTGCTTTTCACTT 3425

Qy   2461 GTGTTTTAAAAATAAGGATTTTTTCTCTTATTTTCTCCCCCTTAGTTTGTGAGAAAACATCTCAA 2520
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3426 GTGTTTTAAAAATAAGGATTTTTTCTCTTATTTTCTCCCCCTTAGTTTGTGAGAAAACATCTCAA 3485

Qy   2521 TAAAGTGCTTTTAAAAAAAAAAAAAAA 2547
          |||||||||||||||  |||||||||
Db   3486 TAAAGTGCTTTCCAAAAAAAAAAAAAAAA 3512
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RESULT 4
US-09-974-592-7
; Sequence 7, Application US/09974592
; Patent No. US20020120121A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: Mackenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Tsang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009004
; CURRENT APPLICATION NUMBER: US/09/974,592
; CURRENT FILING DATE: 2001-10-09
; PRIOR APPLICATION NUMBER: US 09/617,053
; PRIOR FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 08/800,929
; PRIOR FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 7
; LENGTH: 3732
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-974-592-7

Query Match   97.9%; Score 2535.8; DB 10; Length 3732;
Best Local Similarity   99.7%;
Matches 2540; Conservative   0; Mismatches   7; Indels   0; Gaps   0;

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Qy      1 TCTAAGTAGTAGTATCTTGGAAAATTCAGAGAGATACTCATCCTACCTGAATATAAACTGAGAT 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1186 TCTAAGTAGTATCTTGGTAATTCAGAGAGATACTCATCCTACCTACCTGAATATAAACTGAGAT 1245

Qy     61 AAATCCAGTAAAGAAAAGTGTAGTAAATTCTACACATAAGAGAGCTATCTACCTGAATATAAAC 120
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1186 AAATCCAGTAAAGAAAAGTGTAGTAAATTCTACACATAAGAGAGCTATCTACCTGAATATAAAC 1245

Qy    121 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTCATGTGAAGTGTTTAGCTATCAAACAGC 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1246 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTCATGTGAAGTGTTTAGCTATCAAACAGC 1305

Qy    181 ACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAGACTTTCCCAGGTCCCTCGTATC 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1306 ACTGTCACCTACTCATGCACAAAAACTGCCTCCCAAAGACTTTCCCAGGTCCCTCGTATC 1365

Qy    241 AAAACATTAAGAGTATAATGGAAGATAGCACGGATCTTGTCAGATTGGACAAAACAGCAACA 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1366 AAAACATTAAGAGTATAATGGAAGATAGCACGGATCTTGTCAGATTGGACAAAACAGCAACA 1425
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Db   1426 AAAACATTAAGAGTATAATGGAAGATAGCACGGATCTTGTCAGATTGGACAAAACAGCAACA 1485

Qy    301 AACAAAAAAATGAAGTATGAGACTTTTTCCTGTGTGAAACTTCTACAGAATGTCTACAATTCAACTT 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1486 AACAAAAAAATGAAGTATGAGACTTTTTCCTGTGTGAAACTTCTACAGAATGTCTACAATTCAACTT 1545

Qy    361 TCCCCCGCCGGGGGTGGCCTTGTCTCGAGAAAGGAGTCTTGGCTCGTGGCTGGTTTTTATTATACTG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1546 TCCCCCGCCGGGGGTGGCCTTGTCTCGAGAAAGGAGTCTTGGCTCGTGGCTGGTTTTTATTATACTG 1605

Qy    421 GTGTGAATGACAAGGTCAAAATGCTTCTCTGTTGTGTGGGCCTGATGTCGGGATAACTGAAAACTAG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1606 GTGTGAATGACAAGGTCAAAATGCTTCTCTGTTGTGTGGGCCTGATGTCGGGATAACTGAAAACTAG 1665

Qy    481 GAGACAGTCCTATTCAAAAAGCATAAACAGTATATCCTAGCGTGCTTTTATTCAGAATC 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1666 GAGACAGTCCTATTCAAAAAGCATAAACAGCTATATCCTAGCGTGCTTTTATTCAGAATC 1725

Qy    541 TGGTTTCAGCTAGTCTGGGATCCACCTCTAAGAATACGTCTCCAAATGAGAAAACAGTTTTG 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1726 TGGTTTCAGCTAGTCTGGGATCCACCTCTAAGAATACGTCTCCAAATGAGAAAACAGTTTTG 1785

Qy    601 CACATTCATTATATCTCCCACACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1786 CACATTCATTATATCTCCCACACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC 1845

Qy    661 TTTTCTCCCAAACCCTCTTTAATTCTAGAGCAGTTGAAGACATCTCTCATCGAGAGACTAACC 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1846 TTTTCTCCCAAACCCTCTTTAATTCTAGAGCAGTTGAAGACATCTCTCATCGAGAGACTAACC 1905

Qy    721 CCTACAGTTAATGCAAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCCATATGTGGCCAT 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1906 CCTACAGTTAATGCAAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCCATATGTGGCCAT 1965

Qy    781 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTATTATTATAGGAGACCTGGAG 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1966 TAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTATTATTATAGGAGACCTGGAG 2025

Qy    841 ATAGGGTAGCCTGCTTTGCCTGTGGCTGGGAAGCTCAGTAACTGGGAACCAAAGGATGATG 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2026 ATAGGGTAGCCTGCTTTGCCTGTGGCTGGGAAGCTCAGTAACTGGGAACCAAAGGATGATG 2085

Qy    901 CTATGTCAGAACACCCGAGGCCATTTCCAACTGTCCATTTTTGGAAAAATTCTCTAGAA 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2086 CTATGTCAGAACACCCGAGGCCATTTCCAACTGTCCATTTTTGGAAAAATTCTCTAGAA 2145

Qy    961 CTCCTGAGGTTTAGCATTTCAAAATCTGAGCATGCAGACACATGCAGCTCGAATGAGAACAT 1020
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2146 CTCCTGAGGTTTAGCATTTCAAATTCTGAGCATGCAGACACATGCAGCTCGAATGAGAACAT 2205

Qy   1021 TTATGTACTGGCCATCTAGTGTTCCAGTTCAGCCTGAGCAGCTTGCAAGTGCTGGTTTTT 1080
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2206 TTATGTACTGGCCATCTAGTGTTCCAGTTCAGCCTGAGCAGCTTGCAAGTGCTGGTTTTT 2265

Qy   1081 ATTAATGTGGGGTCGCAATGAATGAGTGTCAAATGCTTTTGTTGTGATGGGTTGAGTGTT 1140
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2266 ATTAATGTGGGGTCGCAATGAATGAGTGTCAAATGCTTTTGTTGTGATGGGTTGAGTGTT 2325

Qy   1141 GGGAATCTGGAGATGAGCCATGGGTAGAACATGCCAAGTGTGTTGAGTTCT 1200
          ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2326 GGGAATCTGGAGATGAGCCATGGGTAGAACATGCCAAGTGTGTTGAGTTCT 2385

Qy   1201 TGATAGGAATGAAAGGCCAAGAGTTTGTTTGATGGATTCAAGGTAGATATCCTCATCTTC 1260
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2386 TGATAGGAATGAAAGGCCAAGAGTTTGTTTGATGAGATTCAAGGTAGATATCCTCATCTTC 2445

Qy   1261 TTGAACAGCTGTTGTCAACTTCAGGATACCACTGGAGAAGAGAAATGCTGACCCACCAATTA 1320
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2446 TTGAACAGCTGTTGTCAACTTCAGGATACCACTGGAGAAGAGAAATGCTGACCCACCAATTA 2505

Qy   1321 TTCATTTTGGACCTGGAGAAGTTCTTCAGGAAGATGAATGCTCGTCATGAATGAATACACCCTGTGG 1380
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2506 TTCATTTTGGACCTGGAGAAGTTCTTCAGGAAGATGAATGCTCGTCATGAATGAATACACCCTGTGG 2565
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RESULT 5
US-10-207-655-199
Sequence 199, Application US/10207655
Publication No. US20030118592A1

GENERAL INFORMATION:
- APPLICANT: Ledbetter, Jeffrey A.
- APPLICANT: Hayden-Ledbetter, Martha S.
- TITLE OF INVENTION: BINDING DOMAIN-IMMUNOGLOBULIN FUSION PROTEINS
- FILE REFERENCE: 390069.401C1
- CURRENT APPLICATION NUMBER: US/10/207,655
- CURRENT FILING DATE: 2002-07-25
- NUMBER OF SEQ ID NOS: 426
- SOFTWARE: Patentin version 3.0
- SEQ ID NO 199
- LENGTH: 2531
- TYPE: DNA
- ORGANISM: Homo sapiens

US-10-207-655-199

Query Match 97.8%; Score 2531; DB 14; Length 2531;
Best Local Similarity 100.0%; Pred. No. 0; Mismatches 0;
Matches 2531; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

[This page consists of a paired nucleotide sequence alignment (Query "Qy" vs. Database "Db") spanning the full SEQ ID NO 199 (2531 bp), shown in two column groups with base-position numbers flanking each 60-base alignment row. The alignment lines (vertical bars "|" indicating matched bases) and individual DNA sequences are presented as dense sequence-listing data and are not individually reproduced here.]

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Db   601  CACATTCATTACTCTCCACCTTGGAACATAGTAGTAGCTTGTTCAGTGGTTCTTACTCCAGCC  660
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy

Db   661  TTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGACTAACC  720
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy   661

Db   721  CCTACAGTTATGCAAATGACTACTGAAGAAGAGCCCAGAGTTCTTACCTACCATATGTGGCCAT  780
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy   721  CCTACAGTTATGCAAATGACTACTGAAGAAGAGCCCAGAGTTCTTACCTACCATATGTGGGCCAT

Db   781  TAACTTTTTTGTCACCATCAGAATTGGGCAAGAGCTGGTT  840
          |||||||||||||||||||||||||||||||||||||||||
Qy   781  TAACTTTTTTGTCACCATCAGAATTGGGCAAGAGCTGGAG

Db   841  ATAGGGGTAGCCTTGCTTTTGGCTGTGGTGGGAAGGCTCAGTGAACTCAGGAACCAAAGGATGATG  900
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy   841  ATAGGGGTAGCCTTGCTTTTGCCTGTGTGGTGGGAAGCTCAGTGATAACTGGGAACCAAAGGATGATG

Db   901  CTATGTCAGAACACCCGGAGGCATTTTCCCAACTGTCCCATTTTTTGGAAAATTCTCTAGAAA  960
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy   901  CTATGTCAGAACACCCGGAGGCATTTTCCCAACTGTCCCATTTTTTGGAAAATTCTCTAGAAA

Db   961  CTCTGAGGTTTAGCATTTCAAATCTGACGCAGACACTTGACGGATGAGAACAT  1020
          |||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy   961  CTCTGAGGTTTAGCATTTCAAATCTGACGCAGACAGCATGGACAGCTTCGGAATGAGAACAT

Db  1021  TTATGTGTACTGGGCCATCTAGTGTTCAGTTCAGCGCAGCTTGCAAGTGCTGGTTTTTT  1080
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1021  TTATGTACTGGGCCATCTAGTGTTCAGTTCAGCGCAGCTTGCAAGTGCTGGTTTTTT

Db  1081  ATTAATGTGGGTCGGCCAATGATGATGATGTCAAATGCTTTTTGTTGTGAGGTGGGTTGTT  1140
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1081  ATTAATGTGGGGTCGGCCAATGATGATGTGTCAAATGCTTTTTGTTGTGAGGTGGGTTGTT

Db  1141  GGGAAACTCTGGAGAATGATTCCATGGGTAGAGAACATGCCAAAGTGGGTTCCAAGGTGTGAGTTCT  1200
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1141  GGGAAACTCTGGAGAATGATTCCATGGGTAGAGAACATGCCAAGTGGGTTCCAAGGTGTGAGTTCT

Db  1201  TGATACGGAATGAAAGGCCAAGAGAGTTTGGTTGGTTGATTCAAGGTAGAGTATCCTCATCTTC  1260
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1201  TGATACGGAATGAAAGGCCAAGAGAGTTTGTTGGTTGATTCAAGGTAGAGTATCCTCATCTTC

Db  1261  TTGAACAGCTGTTGTCAACTTCAGATACCAACTGGAGAGAAAATGCTGACCCACCAATTA  1320
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1261  TTGAACAGCTGTTGTCAACTTCAGATACCAACTGGAGAGAAAATGCTGACCCACCAATTA

Db  1321  TTCAATTTGGACCTGGAGAAAGTTCTTCAGAAGATGCGTCATGATGAATAACACCTGTGG  1380
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1321  TTCATTTTGGACCTGGAGAAAGTTCTTCAGAAGATGCGTCATGATGAATAACACCTGTGG

Db  1381  TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGACCTGGTTAAACAAACAGTTCAAAGTA  1440
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1381  TTAAATCTGCCTTGGAAATGGGCTTTAATAGAGACCTGGTTAAACAAACAGTTCAAAGTA

Db  1441  AAATCCTGACAACTGGAGAAGGAAGGAAAACAGTTAATGAGAAAATGGCATCAG  1500
          ||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1441  AAATCCTGACAACTGGAGAAGGAAGGAACTAATAAAACAGTTAATGATGTATATTGTGTCAGCACTTCTAA

Db  1501  ATGCTGAAGATGAAAAAGAGAGGAAGGAAAAACAGACAGCTGAAGAAATGGCATCAG  1560
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1501  ATGCTGAAGATGAAAAAGAGAGGAAGGAAAAACAGACAGCTGAAGAAATGGCATCAG

Db  1561  ATGATTTGTCATTAATTCGGAAGAACAGAATGGCTCTCTCTTCAAACAATTGACACTGTGTC  1620
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1561  ATGATTTGTCATTAATTCGGAAGAACACAGAATGGCTCTCTCTTCAAACAATTGACACATGTGTGC

Db  1621  TTCCTACCCTTGGATAAATCTTTTTAAAGGCCCAATGTAATTAATTATAATAACAGAACATGATATTA  1680
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1621  TTCCTATCCTTGGATAAATCTTTTAAAGGCCCAATGTAATTAATTAATTAATAACAGAACATGATATTA

Db  1681  TTAAACAAAAAACACAGATACCTTTACAAGCGGAGGAGAGAACTGATTGATACCCATTTTGGGTTA  1740
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1681  TTAAACAAAAAAACACAGATAACCTTTTACAAGCGGAGAGGAGAACTGATTGATACCCATTTTGGGTTA
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(right column continued)

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Db  1681  TTAAACAAAAACACAGATACCTTTACAAGCGGAGGAGAGAACTGATTGATACCATTTTGGTTA  1740
Qy

Db  1741  AAGGAAATGCTGCGGGCCCAACATCTTCAAAAAACTGTCTAAAAAGAAAATTGACTCACATTGT  1800
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1741  AAGGAAATGCTGCGGGCCCAACATCTTCAAAAAACTGTCTAAAAAGAAAATTGACTTACATTGTC

Db  1801  ATAAGAAACTTATTTGTGGATAAAGAAATAGAAGTATATTCCAAACAGAAGATAGTTTTCAGGTC  1860
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1801  ATAAGAAACTTAATTTGTGGATAAAGAAATAGAAGTATATTCCAACAGCAGAAGATAGTTTTCAGGTC

Db  1861  TGTCACTGGAGAACAAATTGAGGGAGGTTGCAAGAAAGAACGAACTGGTAAAGTGGTATGG  1920
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1861  TGTCACTGGAGAACAAATTGAGGGAGGTTGCAAGAAGAACGAACTTGTTAAAGTGGTATGG

Db  1921  ACAAAGAAGTTTCTGTTTGGTTGTTATTTATTTATTCCTTGTGGTTCATCTGGTAGTATGCCAGGAAGTGTG  1980
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  1921  ACAAAGAAGTTTCTGTTTGGTTGTTATTTATTTATTCCTTGTGGTTCATCTGGTAGTATGCCAGGAAGTGTG

Db  1981  CCCCTTCTCTAAGAAAAATGCCCTATTTGGCAGGGGTATAATCAAGGGTACTGTTCGTACAT  2040
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Qy  1981  CCCCTTCTCTAAGAAAATGCCCTATTTGGCAGGGGTATAATCAAGGGTACTGTTCGTACAT

Db  2041  TTCTCTCTTAAAGAAAAATAGTCTATATTTTTAACCTGCATAAAAGGTCTTTTAAAATATT  2100
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Qy  2041  TTCTCTCTTAAAGAAAAATAGTCTATATTTTTAACCTGCATAAAAGGTCTTTTAAAATATT

Db  2101  GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGGAAATTATGAGTTTTCTTGAATTAGTAACA  2160
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Qy  2101  GTTGAACACTTGAAGCCATCTAAAGTAAAAAGGGAAATTATGAGTTTTTCAATTAGTAACA

Db  2161  TTCATGTTCTAGTTCTGCTTTGGTACTAATAAAATCTTGTTTTCTGAAAAGATGTATCATATA  2220
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Qy  2161  TTCATGTTCTAGTTCTGCTTTGGTACTAATAAAATCTTGTTTTCTGAAAAGATGTATCATATA

Db  2221  TTTAATCTTAAATCTGTTTAATTTATTTACAAGGGAAGAGATTTAGTGTTGGTGAACTTATTAGTAT  2280
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Qy  2221  TTTAATCTTAAATCTGTTTAATTTATTTACAAGGGAAGAGATTTAGTGTTGGTGAACTTATTAGTAT

Db  2281  GTATGTGTACCTTAAGGGAGTAGTGTCACTGCTTGTTATTGCATCATTCAGTTACTTGG  2340
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Qy  2281  GTATGTGTACCTTAAGGGAGTAGTGTCACTGCTTGTTATTGCATCATTCAGTTACTTGG

Db  2401  CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCGGAATTGTGTGCTTTTCACTT  2460
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Qy  2401  CCAGGAACTCTGGAGTTCATCAGAGTTATGGTGCGGAATTGTGTGCTTTTCACTT

Db  2461  GTGTTTAAAAATAAGGATTTTTCTCTTCTTATTTCCCCCCTAGTTGTTGTGAGAAACATCTCAA  2520
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Qy  2461  GTGTTTAAAAATAAGGATTTTTCTCTTCTTATTTCCCCCCTAGTTTGTTGTGAGAAACATCTCAA

Db  2521  TAAAGTGCTTT  2531
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Qy  2521  TAAAGTGCTTT  2531
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; PRIOR FILING DATE: 2001-05-25
; PRIOR APPLICATION NUMBER: US 60/316,031
; PRIOR FILING DATE: 2001-08-31
; PRIOR APPLICATION NUMBER: US 60/328,403
; PRIOR FILING DATE: 2001-10-12
; PRIOR APPLICATION NUMBER: JP 2001-157043
; PRIOR FILING DATE: 2001-05-25
; PRIOR APPLICATION NUMBER: JP 2001-260681
; PRIOR FILING DATE: 2001-08-30
; PRIOR APPLICATION NUMBER: JP 2001-313175
; PRIOR FILING DATE: 2001-10-10
; NUMBER OF SEQ ID NOS: 488
; SOFTWARE: Patentin Ver. 2.0
;
; SEQ ID NO 337
; LENGTH: 3496
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1160)..(3013)
; US-10-153-668-337

Query Match     97.8%;  Score 2531;  DB 14;  Length 3496;
Best Local Similarity   100.0%;  Pred. No. 0;
Matches 2531;  Conservative 0;  Mismatches 0;  Indels 0;  Gaps 0;
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Qy      1 TCTTAAGTAGTATCTTGGAAATTCAGAGAGATACTCATCCTAYCCTGAAATATAAAACTGAGAT 60
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Db    966 TCTTAAGTAGTATCTTGGAAATTCAGAGAGATACTCATCCTAYCCTGAAATATAAAACTGAGAT 1025

Qy     61 AAAATCCAGTAAAGAAAAGTGTAGTAAATTCTACATAAGAGTCTATCATTGATTTCTTTTGG 120
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Db   1026 AAAATCCAGTAAAGAAAAGTGTAGTAAATTCTACATAAGAGTCTATCATTGATTTCTTTTGG 1085

Qy    121 TGGTAAAAATCTTAGTTCAATGTGAAGAAATTCATGTGAAYTGTTAGTTACTATCAAAACAGC 180
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Db   1086 TGGTAAAAATCTTAGTTCAATGTGAAGAAATTCATGTGAAYTGTTAGTTACTATCAAAACAGC 1145

Qy    181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACACTTTTCCCAGGTCTCCTCGTATC 240
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Db   1146 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACACTTTTCCCAGGTCTCCCTGTATC 1205

Qy    241 AAAACATTAAGAGTATAATGGAAGATAGCACGATCTTGTCGAATTGGACAAACAGCAACA 300
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Db   1206 AAAACATTAAGAGTATAATGGAAGATAGCACGATCTTGTCGAATTGGACAAACAGCAACA 1265

Qy    301 AACAAAAAATGAAGTAYGACTTTTCCTGTGAACTCTACAGAAGAATGCTACATATTCAACTT 360
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Db   1266 AACAAAAAATGAAGTAYGACTTTTCCTGTGAACTCTACAGAAGAATGCTACATATTCAACTT 1325

Qy    361 TCCCCGCCGGGGTGCCCTGTCTCAGAAAGAGTCTTGCTCGTCGTGGTTGTTATATTACTG 420
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Db   1326 TCCCCGCCGGGGTGCCCTGTCTCAGAAAGAGTCTTGCTCGTCGTGGTTGTTATATTACTG 1385

Qy    421 GTGTGAATGACAAGGTCAAATGCTTCTTGTGTTGGCCTGATGAGAAGTGGAAACTAG 480
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Db   1386 GTGTGAATGACAAGGTCAAATGCTTCTTGTGTTGGCCTGATGAGAAGTGGAAACTAG 1445

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Qy    541 TGGTTTCAGCTAGTCTGGGATCCACCCTCTAAGAAATAGGTCTTCAAATGAGAAACAGTTTTG 600
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Db   1506 TGGTTTCAGCTAGTCTGGGATCCACCCTCTAAGAAATAGGTCTTCAAATGAGAAACAGTTTTG 1565

Qy    601 CACATTCATTATCTCCCACCTTGGAACATAGTACTAGTTGTTGTTCAGTGGTTCTTACTCCCAGCC 660
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Db   1566 CACATTCATTATCTCCCACCTTGGAACATAGTACTAGTTGTTGTTCAGTGGTTCTTACTCCCAGCC 1625

Qy    661 TTTCTCCAAACCCTCTTAATTCTAGAGAGCAGTTGAGAGACAATCXYCTTCATCGAGAGGACTAACC 720
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Qy    721 CCTACAGTTATGCAAATGAGTGAGTACTGAAGAGAAGCCCAGATTTCTTACCTACCATATGTGGCCAT 780
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Db   1686 CCTACAGTTATGCAAATGAGTGAGTACTGAAGAGAAGCCCAGATTTCTTACCTACCATATGTGGCCAT 1745

Qy    781 TAAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATTAGGACTTGGAG 840
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Db   1746 TAAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATTAGGACTTGGAG 1805

Qy    841 ATAGGGGTAGCCTCGTTTGCCTGTGGTGGGAAGCTCAGTAACTAACGAACCAAAGATGATG 900
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Db   1806 ATAGGGGTAGCCTCGTTTGCCTGTGGTGGGAAGCTCAGTAACTAACGAACCAAAGATGATG 1865

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Qy   1081 ATTATGTGGGTCGGCAATGATGATGTCAAATGCTTTTGTTGTTGTCGATGGTGGCTTGAGGTGTT 1140
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Db   2046 ATTATGTGGGTCGGCAATGATGATGTCAAATGCTTTTGTTGTTGTCGATGGTGGCTTGAGGTGTT 2105

Qy   1141 GGGAAATCTGGAGATGATCCATGGGTAGAAGTGCCAAGTGGTTTTCCAAGGTGTGAGTTCT 1200
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Db   2106 GGGAAATCTGGAGATGATCCATGGGTAGAAGTGCCAAGTGGTTTTCCAAGGTGTGAGTTCT 2165

Qy   1201 TGATACGGAATGAAAGGCCAAGAGAGTTTGTTGATGAAGTAGAGATATCCTCATCTTC 1260
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Db   2166 TGATACGGAATGAAAGGCCAAGAGAGTTTGTTGATGAAGTAGAGATATCCTCATCTTC 2225

Qy   1261 TTGAACAGCTGTTGTCAACTTCAGATACCACTGAGAAGAAAATGCTTGACCCACCACAATTA 1320
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Qy   1321 TTCATTTGGACCTGGAGAAGTTCTTCAGAAGATGCTGTCATGATGAATACACTGTGG 1380
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Db   2286 TTCATTTGGACCTGGAGAAGTTCTTCAGAAGATGCTGTCATGATGAATACACTGTGG 2345

Qy   1381 TTAAAATCTGCCTTGGAAATGGGGCTTTAATTAGAGAGACGACCTGGTGAAAACAACAGTTCAAAGTA 1440
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Db   2346 TTAAAATCTGCCTTGGAAATGGGGCTTTAATTAGAGAGACGACCTGGTGAAAACAACAGTTCAAAGTA 2405

Qy   1441 AAATCCTGACAAACTGGAGAGAACTATTAAAACAGTTAAGATAGTGTCGCACACTTCTAA 1500
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Db   2406 AAATCCTGACAAACTGGAGAGAACTATTAAAACAGTTAAGATAGTGTCGCACACTTCTAA 2465

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Db   2466 ATGCTGAAGATGAAAAAAGAGAGAGGAGGAGGAGAAAACAAGCTGAAGAAAATGGCATCAG 2525

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Db   2526 ATGATTTGTCATTAATTCGGAAGAGAAACAGAATGGCTCTCTTCAACAATTGACATGTGC 2585

Qy   1621 TTCCTATCCTGGATAATCTTTTAAAGGCCAAATGTAATTAAAAACAGGGAGAACATGATATTA 1680
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Qy   1681 TTAAACAAAAAAACACAGATACCTTTACAAAGCGAGAGAACTGATTGATACCATTTTGGTTA 1740
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RESULT 7
US-09-201-936-7
; Sequence 7, Application US/09201936
; Publication No. US20020187946A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: MacKenzie, Alexander E.
; APPLICANT: Baird, Stephen
; APPLICANT: Liston, Peter
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; FILE REFERENCE: 07891/003003
; CURRENT APPLICATION NUMBER: US/09/201,936
; CURRENT FILING DATE: 1998-12-01
; EARLIER APPLICATION NUMBER: 09/011,356
; EARLIER FILING DATE: 1998-02-04
; EARLIER APPLICATION NUMBER: PCT/IB96/01022
; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 08/576,956
; EARLIER FILING DATE: 1995-12-22
; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 7
; LENGTH: 2580
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: variation
; LOCATION: (2412)...(2412)
; OTHER INFORMATION: N may be any nucleotide
; US-09-201-936-7

Query Match   97.1%; Score 2514.8; DB 10; Length 2580;
Best Local Similarity   99.4%; Pred. No. 0;
Matches 2522; Conservative   0; Mismatches 14; Indels   0; Gaps   0;

Qy   1 TCTAAGTAGTATCTTGAAAATTCAGAGAGAATACTCATCCTACCTGAATATAAAACTGAGAT 60
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Qy   121 TGGTAAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAAATGTTTAGCTATCAAAACAGC 180
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Db   164 TGGTGAAAATCTTAGTTCATGTGAAGAAAATTTCATGTGAAATGTTTAGCTATCAAAACAGT 223

Qy   181 ACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTTCCCTCGTATC 240
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Qy   241 AAAACATTAAGAGTATTAATGGAAGATAGCACAGATCTTGTCAGATTGGACAAACAGCAACA 300
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Db   284 AAAACATTAAGAGTATTAATGGAAGATAGCACAGATCTTGTCAGATTGGACAAACAGCAACA 343

Qy   301 AACAAAAAATGAAGTATGACTTTTCCGTGGAACTCTACAGAAGTCTACACATATTCAACTT 360
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Qy   421 GTGTGAAATGACAAAGTCAAATGCTTCTCGTTGGTGGCCTGATGCTGGAATAACTGGAAACTAG 480
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Db   464 GTGTGAAATGACAAAGTCAAATGCTTCTCGTTGTGGCCTGATGCTGGAATAACTGGAAACTAG 523

Qy   481 GAGACAGTCCTATTCAAAAGCATAAAACAGCTATATCCTAGCTGTAGCTTTATTCAGAATC 540
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Qy   541 TGGTTTCAGCTAGTCTGGGATCCACCCTCTAAGAATACGTCTCCAATGAGAGAAAACAGTTTTG 600
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Qy   601 CACATTCATTATCTCCCACCTTGGAACATAGTACTTTGTTCAGTGTTCTTACTCCAGCC 660
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Db   644 CACATTCATTATCTCCCACCTTGGAACATAGTACTTTGTTCAGTGTTCTTACTCCAGCC 703

Qy   661 TTTCTCCAAACCCTCTTAAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGACTAACC 720
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Qy   721 CCTACAGTTATGCAATGATACTGAAGAAGCCAGATTTTCTTACCATATATGTGGCCAT 780
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Qy   1801 ATAAGAACTTATTTGTGGAATAAGAAATATGAAGTATATATTCCAACAGAAGAAGATGTTTCAGGTC 1860
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Db   2766 ATAAGAACTTATTTGTGGATAAGAAATATGAAGTATATTCCAACAGAAGAAGATGTTTCAGGTC 2825

Qy   1861 TGTCACTGGAAGAACAATTGAGGAGGTTGCAAGAAGAAC(GAACTTGTAAAGTGTGTATGG 1920
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Qy   1921 ACAAAGAAGTTTCTGTTGGTATTTATTCCTTGGTGGTCATCCAGGAAATGTG 1980
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Db   2886 ACAAAGAAGTTTCTGTTGGTATTTATTCCTTGGTGGTCATCCAGGAAATGTG 2945

Qy   1981 CCCCTTCTCTAAGAAAAATGCCCTATTTGCAGGGGGTATAATCAAGGGTACTGTTTCGTACAT 2040
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Qy   2041 TTCTCTCTTAAAGAAAAATTAGTCTATATATTTTAAACCTGCATAAAAAAAGGTCTTTAAAATATT 2100
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Db   3126 TTCATGTTCTAGTCTGCTTGGGTACTAATAATAATCTTGTTTCTGAAAAAGATGGTATCATATATA 3185

Qy   2221 TTTAAATCTTAATCTGTTTATTTACAAGGGAGGATTTATTGTTTATGTTGAACTTATTAGTAT 2280
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Db   3186 TTTAAATCTTAATCTGTTTATTTACAAGGGAGGATTTATTGTTTATGTTGAACTTATTAGTAT 3245

Qy   2281 GTATGTGTTACCTAAGGGAGGTGTGCACTGCTTGTTGTTATGCATCATTCAGGAGTTACTGG 3305
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Qy   2401 CCAGGAACTCTCGGAGTTCATCAGAGGTTAATGGTGGCCGAATTGTCTTGGTGCTTTTCACTT 2460
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Db   3366 CCAGGAACTCTCGGAGTTCATCAGAGTTATGTGCCGAAATTGTCTTGGTGCTTTTCACTT 3425

Qy   2461 GTGTTTTAAAATAAGGATTTTTCTCTCTATTTCTCCCCCTAGTTTGTTGTGAGAAAAACATCTCAA 2520
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Db   3426 GTGTTTTAAAATAAGGATTTTTCTCTCTATTTCTCCCCCTAGTTTGTTGTGAGAAAAACATCTCAA 3485

Qy   2521 TAAAAGTGCTTTT 2531
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Db   3486 TAAAAGTGCTTTT 3496

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Qy    841  ATAGGGTAGCCTTGCTTTGCCTGTTGGTTGGGAAGCTCAGTATA.CTGGGAAGGATGATG  900
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Db    884  ATAGGGGTAGCCTGCTTGCCTGTGGTTGGGAAGCTCAGTAACCAAAGGATGATG       943

Qy    901  CTATGTCAGAACACCGGAGGCATTTTCCCACTGTCCATTTTTGGAAAATTCTCTAGAAA   960
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Db    944  CTATGTCAGAACACCGGAGGCATTTTCCCAACTGTCC.ATTTTTGGAAAATTCTCTAGAAA 1003

Qy    961  CTCTGAGGTTTAGCATTTCAAATCTGAGCATGCAGCAGCTCGAATGAGAAACAT       1020
           |||||||||||||||||||||||||||||||||||||||||||||||||
Db   1004  CTCTGAGGTTTAGCATTTCAAATCTGAGCATGCAGCAGCTCGAATGAGAAACAT       1063

Qy   1021  TTATGTACTGGGCCATCTAGTGTTTCAGTTCAGCCTGAGCAGCTTGCAAGTGCTCGTTT   1080
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1064  TTATGTACTGGGCCATCTAGTGTTTCAGTTCAGCCTGAGCAGCTTGCAAGTGCTCGTTTTT 1123

Qy   1081  ATTAGTGGGGTCGCAATGATGAGTCAAAGCTTTTGTTGTTGTGATGGGTGGCTTGAGGTGTT 1140
           ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1124  ATTAGTGGGGTCGCAATGATGAGTCAAAAGCTTTTGTTGTGATGGGTGGCTTGAGGTGTT   1183

Qy   1141  GGGAATCTGGAGATGAATCCATGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 1200
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1184  GGGAATCTGGAGATGAATCCATGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGAGTTCT 1243

Qy   1201  TGATACGGAATGAAAGGCCAAGAGTTTGTTGATGATGAGATATCCTCAATCTTC       1260
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1244  TGATACGGAATGAAAGGCCAAGAGTTTGTTGATGATGAGATATCCTCAATCTTC       1303

Qy   1261  TTGAACAGCTGTGTTGTCAACTTCAGATACCACTGGGAGAGAGAAATGCTGACCCACCAAATTA 1320
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1304  TTGAACAGCTGTGTTGTCAACTTCAGATACCACTGGGAGAGAGAAATGCTGACCCACCAAATTA 1363

Qy   1321  TTCATTTTGGGACCTGGAGAAAGTTCTTCAGAGAGAGCTGCATGATGAATACACCTGTGG   1380
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1364  TTCATTTTGGGACCTGGAGAAAGTTCTTCAGAGAGAGCTGCATGATGAATACACCTGTGG   1423

Qy   1381  TTAAAATCTGCCTTGGAAAATGGGCTTTAATAGAGACCTGGTGAAACAGTTCAAAGTA    1440
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1424  TTAAAATCTGCCTTGGAAAATGGGCTTTAATAGAGACCTGGTGAAACAGTTCAAAGTA    1483

Qy   1441  AAAATCCTGACAACTGGAGAGAACTATAAAAACAGTTAATGATAATTGTCAGCACTTCTTA   1500
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1484  AAATCCTGACAACTGGAGAGAACTATAAAAACAGTTAATGATAATTGTCAGCACTTCTTA   1543

Qy   1501  ATGCTGAAGATGAAAAGAGAGGAGGAAAACAAGCTGAAGAAAATGGCATCAG       1560
           ||||||||||||||||||||||||||||||||||||||||||||||||
Db   1544  ATGCTGAAGATGAAAAGAGAGGAGGAAAACAAGCTGAAGAAAATGGCATCAG       1603

Qy   1561  ATGATTTGTCATTPAATTCGGAAGAACAGAATGGCTCTCCTTTCAACAATTGACATGTGTGC 1620
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1604  ATGATTTGTCATTPAATTCGGAAGAACAGAATGGCTCTCCTTTCAACAATTGACATGTGTGC 1663

Qy   1621  TTCCTATCCTGGATAATCTTTTTAAAGGCCAAATGTAATTAATAAAACAGGAACAGATGATATTA 1680
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1664  TTCCTATCCTGGATATATCTTTTTAAAGGCCAAATGTAATTAATAAACAGGAACAGATGATATTA 1723

Qy   1681  TTAAACAAAAAACACAGATACCTTTACAAGCGGAGGAACTGATTGATACCATTTTGGTTA  1740
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1724  TTAAACAAAAAACACAGATACCTTTACAAGCGGAGGAACTGATTGATACCATTTTGGTTA  1783

Qy   1741  AAGGAAAATGCTGCGGCCCACATCTTCTAAAAAAGAAATTGACTCTACATTGT        1800
           ||||||||||||||||||||||||||||||||||||||||||||||||
Db   1784  AAGGAAAATGCTGCGGCCCACATCTTCTAAAAAAACTGTCTAAAAAAGAAATTGACTCTACATTGT 1843

Qy   1801  ATAAGAACTTATTTGTGGATAAGAATATGAAGTATATATTCCAACAGAGAAGATGTTTCAGGTC  1860
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1844  ATAAGAACTTATTTGTGGATAAGAATATGAAGTATATTCCAACAGAGAAGATGTTTCAGGTC  1903

Qy   1861  TGTCACTGGAAGAACAATTGAGGAGGTTGCAAGAAGAACTTGTGTAAAGTGTGTATGG    1920
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1904  TGTCACTGGAAGAACAATTGAGGAGGTTGCAAGAAGAACGAACTTGTGTAAAGTGTGTATGG  1963

Qy   1921  ACAAAGAAGTTTCTTGTTATTTTATTCCTTGTGTGGTCATCTGGTAGTATGCCAGGAAGTG  1980
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1964  ACAAAGAAGTTTCTTGTTGTTTATTTATTCCTTGTGTGGTCATCTGGTAGTATGCCAGGAAGTGTG 2023

Qy   1981  CCCCTTCTCTAAGAAAAAAGCCCTATTGTTGCAAGGGGGTATATAAGGGGTACTGTTGTTGGTACAT 2040
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2024  CCCCTTCTCTAAGAAAAATGCCCCTATTTGCAGGGGGTATAATCAAGGGGTACGTTCGTACAT 2083

Qy   2041  TTCTCTCTTAAAGAAAATAGTCTATATATTTTAACCTGACATAAAAAAGGTCTTTAAAATATT 2100
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2084  TTCTCTCTTAAAGAAAATAGTCTATATATTTTAACCTGACATAAAAAAGGTCTTTAAAATATT 2143

Qy   2101  GTTGAACACTTGAAGCCATCTAAAGTAAAAAAGAGGGAATTATGAGTTTTTTCAAGTTAACA 2160
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2144  GTTGAACACTTGAAGCCATCTAAAGTAAAAAAGAGGGAATTATGAGTTTTTTCAAGTTAACA 2203

Qy   2161  TTCATGTTCTAGTCTGCTTGGTACTAATAAATCTGTTTCTGAAAAGAATGGTATCATATA  2220
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2204  TTCATGTTCTAGTCTGCTTGGTACTAATAAATCTGTTTCTGAAAAGAATGGTATCATATA  2263

Qy   2221  TTTAATCTTAAATCTGTTTATTTACAAAGGGAAGAGATTTATGTTTTGGGTGAAACTATTAGTTAT 2280
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2264  TTTAATCTTAAATCTGTTTATTTACAAAGGGAAGAGATTTATGTTTTGGGTGAAACTATTAGTTAT 2323

Qy   2281  GTATGTGTACCTAAGGGAGTAGTGTCACTGCTTGTTGTTATGGCATCATTCAGGAGTTACTGG 2340
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2324  GTATGTGTACCTAAGGGAGTAGCGTCNCTGCCTTGTTGTTATGCATCATTTCAGGAGTTACTGG 2383

Qy   2341  ATTTGTTGTTCTTTCAGAAAAGCTTTGAATACTAATTATTAGTGTAGAAAAAGAACTGGAAA 2400
           ||||||||||||||||||| |||||||||||||||||||||||||||
Db   2384  ATTTGTTGTTCTTTCAGAAAAGCTTTGAAAAACTAAATTATAGTGTAGAAAAGAGAACTGGAAA 2443

Qy   2401  CCAGGAAACTCTGGAGTTCATCAGAGTTATAGGGTGCGGAAATGTGTCTTTGGCTTTTGCACTT 2460
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2444  CCAGGAAACTCTGGAGTTCATCAGAGTTATGGTGCCGGAAATGTGTCTTTGGCTTTTGCACTT 2503

Qy   2461  GTGTTTTAAAATAAGGATTTTTCTCTTATTTCTCCCCCTAGTTTGTTGTGAGAAAACATCTCAA 2520
           ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2504  GTGTTTTAAAATAAGGATTTTTCTCCTTATTTCTCCCCCTAGTTTGTTGTGAGAAAACATCTCAA 2563

Qy   2521  TAAAGTGCTTTAAAAA    2536
           ||||||||||||||||
Db   2564  TAAAGTGCTTTAAAAA    2579
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; ATTORNEY/AGENT INFORMATION:
;   NAME: FERRARO, GREGORY D.
;   REGISTRATION NUMBER: 36,134
;   REFERENCE/DOCKET NUMBER: 325800-387
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE: 201-994-1700
;     TELEFAX: 201-994-1744
; INFORMATION FOR SEQ ID NO: 1:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 1435 BASE PAIRS
;     TYPE: NUCLEIC ACID
;     STRANDEDNESS: SINGLE
;     TOPOLOGY: LINEAR
;     MOLECULE TYPE: cDNA
US-08-464-588-1

Query Match             55.2%;  Score 1430.2;  DB 8;  Length 1435;
Best Local Similarity   99.8%;  Pred. No. 2.5e-291;
Matches 1432;  Conservative 0;  Mismatches 3;  Indels 0;  Gaps 0;
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Qy  726 AGTTATGCAAATGAGTACTGAAGAAGGCCAGAGATTTCTTACCTACCATAGTGTGGCCATTAACT 785
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Db    1 AGTTATGCAAATGAGTACTGAAGAAGGCCAGAGATTTCTTACCTACCATAGTGTGGCCATTAACC 60

Qy  786 TTTTTGTCACCATCAGAATTGGGCAAGAGAGCTGGTTTTTATTTATATAGGAGACCTGGAGATAGG 845
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 TTTTTGTCACCATCAGAATTGGGCAAGAGAGCTGGTTTTTATTTATATAGGAGACCTGGAGATAGG 120

Qy  846 GTAGCCTGCTTTGCCTGTGGGGAGGAAGCTCAGTGAACTCGGAACCAAAGGATGATGCTATG 905
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  121 GTAGCCTGCTTTGCCTGTGGGGAGGAAGCTCAGTGAACTCGGAACCAAAGGATGATGCTATG 180

Qy  906 TCAGAAACACCGGGAGCATTTTCCCAACTGTCCATTTTTTGGAAAATTCTCTAGAAACTCTG 965
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  181 TCAGAAACACCGGGAGCATTTTCCCAACTGTCCATTTTTTGGAAAATTCTCTAGAAACTCTG 240

Qy  966 AGGTTTAGCATTTCAAATCTGAGCAATGCAGCAACACATGCGAAATGAGGAACAATTTATG 1025
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  241 AGGTTTAGCATTTCAAATCTGAGCAATGCAGCAACACATGCGAAATGAGGAACAATTTATG 300

Qy 1026 TACTGGGCCATCTAGTTGTTCCAGTTCAGCCTGAGCAGCTTGCAAGTGCTGGTTTTTATTAT 1085
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  301 TACTGGGCCATCTAGTTGTTCCAGTTCAGCCTGAGCAGCTTGCAAGTGCTGGTTTTTATTAT 360

Qy 1086 GTGGGTCGCAATGATGATGTCAAATGCTTTTGTTGTGTGATGGTGGCTTGAGGTGTTGGGAA 1145
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  361 GTGGGTCGCAATGATGATGTCAAATGCTTTTGTTGTGTGATGGTGGCTTGAGGTGTTGGGAA 420

Qy 1146 TCTGGAGATGATCCATGGGGTAGAACATGCCAAGTGTGCCTTGAGGTTGCTTGAGTTCTTGATA 1205
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  421 TCTGGAGATGATCCATGGGGTAGAACATGCCAAGTGTGCCTTGAGGTTGCTTGAGTTCTTGATA 480

Qy 1206 CGAATGAAAGGCCAAGAGTTTGTTGATGAGATTCAAGGTAGATATCCTCATCTTCTTGAA 1265
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  481 CGAATGAAAGGCCAAGAGTTTGTTGATGAGATTCAAGGTAGATATCCTCATCTTCTTGAA 540

Qy 1266 CAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCCACCACCAATTATTCAT 1325
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  541 CAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCCACCACCAATTATTCAT 600

Qy 1326 TTTGGACCTGGAGAAAGTTCTTCAGAAGATGCTGTCATGATGAATACACCTGTGGTTTAAA 1385
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  601 TTTGGACCTGGAGAAAGTTCTTCAGAAGATGCTGTCATGATGAATACACCTGTGGTTTAAA 660

Qy 1386 TCTTGCCTTGGAAATGGGCTTTTAATAGAGACCTGGTGAAACAAACAGTTCAAAGTAAAATC 1445
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  661 TCTTGCCTTGGAAATGGGCTTTTAATAGAGACCTGGTGAAACAAACAGTTCAAAGTAAAATC 720

Qy 1446 CTGACAACTGGAGAGAACTATAAAACAGTTAATGATATTGTGTCAGCCACTTCTTAAATGCT 1505
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  721 CTGACAACTGGAGAGAACTATAAAACAGTTAATGATATTGTGTCAGCCACTTCTTAAATGCT 780

Qy 1506 GAAGATGAAAAAGAGGAGGAAGAACAAGCTGAAGAAAATGGCATCAGATGAT 1565
        ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  781 GAAGATGAAAAAGAGGAGGAAGAACAAGCTGAAGAAAATGGCATCAGATGAT 840
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RESULT 9
US-10-323-643-1

; Sequence 1, Application US/10323643
; Publication No. US20030108552A1
; GENERAL INFORMATION:
;   APPLICANT: He, et al.
;   TITLE OF INVENTION: Human Inhibitor of Apotosis Gene 1
;   FILE REFERENCE: PF165P1D1
;   CURRENT APPLICATION NUMBER: US/10/323,643
;   CURRENT FILING DATE: 2002-12-20
;   PRIOR APPLICATION NUMBER: 08/464,588
;   PRIOR FILING DATE: 1995-06-05
;   PRIOR APPLICATION NUMBER: PCT/US95/05922
;   PRIOR FILING DATE: 1995-05-11
;   NUMBER OF SEQ ID NOS: 10
;   SOFTWARE: Patentin version 3.1
; SEQ ID NO 1
;   LENGTH: 1435
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;     NAME/KEY: CDS
;     LOCATION: (10)..(1326)
;   OTHER INFORMATION:
US-10-323-643-1

Query Match             55.2%;  Score 1430.2;  DB 14;  Length 1435;
Best Local Similarity   99.8%;  Pred. No. 2.5e-291;
Matches 1432;  Conservative 0;  Mismatches 3;  Indels 0;  Gaps 0;
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Qy   726  AGTTATGCAATGAGTACTGAAGAAGCCAGATTTTTCTTACCTTACCCATATGTGGCCATTAAACT  785
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     1  AGTTATGCAATGAGTACTGAAGAAGCCAGATTTTCTTACCTTACCCATATGTGGCCATTAAACC   60

Qy   786  TTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATAGGAGACCTGGAGATAGG  845
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    61  TTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATAGGAGACCTGGAGATAGG  120

Qy   846  GTACCCTGCTTTGCCTGTGGTGGGAAGTCAGTAACTGGAACCAAAGGAGATGGATGCTATG  905
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121  GTACCCTGCTTTGCCTGTGGTGGGAAGTCAGTAACTGGTGCCAAAGGAGATGGATGCTATG  180

Qy   906  TCAGAACACCGGAGGCCATTTCCCAACTGTCATTTGGAAAATTCTCTAGAAACTCTG  965
          |||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181  TCAGAACACCGGAGGGCCATTTCCCAACTGTCCATTTTGGAAAATTCTCTAGAAACTCTG  240

Qy   966  AGGTTTAGCATTTCAAATCTGAGCATGCAGCTCGGATGAGGAACATTTATG  1025
          |||||||||||||||||||||||||||||||||||||||||||
Db   241  AGGTTTAGCATTTCAAATCTGAGCATGCAGCTCGGATGAGGAACATTTATG  300

Qy  1026  TACTGGCCATCTAGTGTTCCAGTTCAGCCTGAGCAGCTTGCCAAGTGACCTGGTTTTTATTAT  1085
          ||||||||||||||||||||||||||||||||||||||||||||||
Db   301  TACTGGCCATCTAGTGTTCCAGTTCAGCCTGAGCAGCTTGCCAAGTGACTGGTTTTTATTAT  360

Qy  1086  GTGGGTCGCAATGATGATGGTGTCAAANGCTTTTGTTGTTGTGTGAGGTGGGCTTGAGGTGTGGGAA  1145
          |||||||||||||||||||||||||||||||||||||||||||||||
Db   361  GTGGGTCGCAATGATGATGTCAAANTGCTTTTGTTGTTGTGTGGCTTGGCTTGAGGTGTTGGGAA  420

Qy  1146  TCTGGAGATGATCCATGGGGTAGAACATGCCAAGTGGGTTTTCCAAGGTGTGAGTTCTTGATA  1205
          ||||||||||||||||||||||||||||||||||||||||||||
Db   421  TCTGGAGATGATCCATGGGGTAGAACATGCCAAGTGGGTTTTCCAAGGTGTGAGTTCTTGATA  480

Qy  1206  CGAATGAAAGGCCAAGAGTTTGTTGTTGGAATTCAAGGTAGATATCCTCATCTTCTTGAA  1265
          ||||||||||||||||||||||||||||||||||||||||||||
Db   481  CGAATGAAAGGCCAAGAGTTTGTTGTTGAATTCAAGGTAGATATCCTCATCTTCTTGAA  540

Qy  1266  CAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCCACCAAATATTCAT  1325
          ||||||||||||||||||||||||||||||||||||||||||||
Db   541  CAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCCACCAAATATTCAT  600

Qy  1326  TTTGGAACCTGGGAAAATGGGGCTTTTAATAGAGAACACTGTGGTTAAA  1385
          ||||||||||||||||||||||||||||||||||||||
Db   601  TTTGGACCTGGGAAAGTTCTTCAGAAGATGCTGTCATGANGAATACACCTGTGGTTAAA  660

Qy  1386  TCTGCCTGGAAATGGGCTTTAATAGACACCTGGGTGAAAACAACAGTTCAAAGTAAAATC  1445
          |||||||||||||||||||||||||||||||||||||||||||
Db   661  TCTGCCTGGAAATGGGCTTTTAATAGACACCTGGGTGAAAACAACAGTTCAAAGTAAAATC  720

Qy  1446  CTGACAACTGGAGAACTATAAAACAGTTAATGATAATTGTGTCAGCACTTCTAAATGCT  1505
          |||||||||||||||||||||||||||||||||||||||||
Db   721  CTGACAACTGGAGAACTATAAAACAGTTAATGATAATTGTGTCAGCACTTCTTAAATGCT  780

Qy  1506  GAAGATGAAAAAAGAGAGGAGGAGGAAAAAACAACCTGAGAAAATGCATCAGATGAT  1565
          ||||||||||||| ||||||||||||||||||||||||||
Db   781  GAAGATGAAAAAGAAGGAGGAGGAGGAAAAAACAACCTGAGAGAAATGGCATCAGATGAT  840

Qy  1566  TTGTCATTAATTCGGAAGAACAAAATGGCCTCTCTTTTCAACAATTGACATGGTGGTTCCT  1625
          |||||||||||||||||||||||||||||||||||||||||||||||
Db   841  TTGTCATTAATTCGGAAGAACAGAGAACATGGCTCTCTTTTCAACAATTGACATGTGTCTTCCT  900

Qy  1626  ATCCTGGATAATCTTTTAAAGGCCAATGTAATTAATAAACAGAGAACAATGATATATTATTAAA  1685
          ||||||||||||||||||||||||||||||||||||||||||||
Db   901  ATCCTGGATAATCTTTTTAAAGGCCAATGTAATGATAATGATATATTATTAAA  960

Qy  1686  CAAAAAACACAGATACCTTTACAACGGAGGAGAACTGATTGATACCATTTGGTTAAAGGA  1745
          |||||||||||||||||||||||||||||||||||||||||
Db   961  CAAAAAACACAGATACCTTTACAACGGAGGAGAACTGATTGATACCATTTGGTTAAAGGA  1020

Qy  1746  AATGCTGCGGCCAACATCTTCAAAAAAACTGTCTTAAAAAGAAAATTGACTCTACACTTGTATAAG  1805
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Db  1021  AATGCTGGGCCAACATCTTCAAAAAACTGTCTTCTACACTTTTGTATATAG  1080
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Qy  1806  AACTTATTGTTGGATAAGAGAATATGAAGTATTAATTCCAACAGAGAACAGTGTTTCAGGTCTGTCA  1865
          |||||||||||||||||||||||||||||||||||||||||||||
Db  1081  AACTTATTGTTGGATAAGAGAATATGAAGTATTAATTCCAACAGAGAACAGTGTTTCAGGTCTGTCA  1140

Qy  1866  CTGGAAGAACAAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTTGTATGGACAAA  1925
          ||||||||||||||||||||||||||||||||||||||||||||
Db  1141  CTGGAAGAACAAATTGAGGAGGTTGCAAGAAGAACGAACTTGTAAAGTGTTGTATGGACAAA  1200

Qy  1926  GAAGTTCTGTTGTATTTAATTCCTTGTGTCATCTGGTTAGTAGTATGCCAGAAATGTGCCCCT  1985
          |||||||||||||||||||||||||||||||||||||||||||||
Db  1201  GAAGTTCTGTTGTATTTAATTCCTTGTGTCATCTGGTTAGTAGTATGCCAGAAATGTGCCCCT  1260

Qy  1986  TCTCTAAGAAAATGCCCTATTTGCAGGGGTATAATCAAGGGGTGTACTGTTCGTACATTTCTC  2045
          |||||||||||||||||||||||||||||||||||||||||||||
Db  1261  TCTCTAAGAAAATGCCCTATTTGCAGGGGTATAATCAAGGGGTGTACTGTTCGTACATTTCTC  1320

Qy  2046  TCTTAAAGAAAAAATAGTTCTATATTTAAACCTGCATAAAAAAAGGTGTCTTTAAAATATTGTTGA  2105
          ||||||||||||||||||||||||||||||||||||||||||
Db  1321  TCTTAAAGAAAAATAGTCTATATTTAAACCTGCATAAAAAGGTCTTTAAAATATTGTTGA  1380

Qy  2106  ACACTTGAAGCCATCTAAAGTAAAAGGAAATTATGAGTTTTTCAATTAGTAGTAACA  2160
          |||||||||||||||||||||||||||||||||||||||||
Db  1381  ACACTTGAAGCCATCTAAAGTAAAAGGAAATTATGAGTTTTTCAATTAGTAGTAACA  1435
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Qy   140  TGTGAAGAAATTTCATGTGAAGTGTTTAGCTATCAAACAGCACTGTCACTTACTCATGCA   199
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Db   420  TGGTGAGAGAACTTCATCTGGAAGTTTAAGCGGTCAGAAATACTATT--ACTACTCATGGA   477

Qy   200  CAAAACTCCCTCCCAAAGACTTTTCCCAGGTTCCCTCGTATCAAAAACATTAAGAGTATAAT   259
               ||||| || |||| |||||||  || ||| ||  |  | || ||||  || || |||| |||
Db   478  CAAAACTGTCTCCCAGAGACTTCGGCCAAAGGTTACCTTACCAAAAAAACTTAAACGTATAAT   537

Qy   260  GGAAGATAGTCACGGATTCTTGTTCAGATTGGACAAAACAGCAACAAAACAAAAAAATGAAGTATGA   319
             |||| || || || ||||||||||||| ||||||||  |  || |  |||| |||||||| ||
Db   538  GGAGGAGAGACAGAATCTTGTTCAAAATTGGACGGAGGAGGAGGGAAGAAAAAATGAAGTTTGA   597

Qy   320  CTTTTTCCTGTGAACTCTACAGAAATTGTTCTACATATTTCCCGCGGGGTTGCCTGTT   379
            ||||| ||| || | | ||| ||||  |  | ||  |  |||| | ||||| |||| ||
Db   598  GTGCTTCTGCTGTGGGCCCTGATTGTGGATAACTGGAAAACAGTGCCTGTTGCCAC   657

Qy   380  CTCAGAAAAGGAGTCTTGCTCGTGCTTGGTGTTTTTTTTAAATACTGGTGTGAATGACAAGGTCAA   439
             || ||||  ||||  || |  || ||| ||||| |||| | || |||| || |||| || ||
Db   658  CTCAGAGAGGGAGTCCTCGGCTCGGCTGCCTGTTTTTTATTACAGTGTGTGAATGACAAAGTCAA   717

Qy   440  ATGCTTCTGTTGTGTGGCCCTGATGCTGGAAACTAGGAGACAGTGCCTATTCAAAA   499
               |||| |  || ||| | || |||| | | | || || |||||  | |  |  ||||
Db   718  GTGCTTCTGCTGTGGGCCCTGATTGTGGATAACAGTGCCTGTTGCCAC   777

Qy   500  GCATAAACAGCTATATCCTTAGCTGTGTAGCTTTATTCAGAATCTGGTTTCAGCTAGTCTGGG   559
              |||| | || |  |  ||   |||||| ||  ||| |  |||| |||||||||||||||||||
Db   778  GCACAGACAGTTCTATCCCAGCTGCAGCTTTGTTACGAACTCTGCTTTGCAGCCCAGTCTGCA   837

Qy   560  ATCCACCTTCTAAGAAATACGTTCTCCAAATGAGAAACAGTTTTTGCACATTCATTATCTCCCAC   619
            || || || ||| ||||| || |  | |||| ||||  |||| | ||||| ||  | |||| ||
Db   838  GTCTCCATCTAAGAATATGTCTCCTGTGAAAAGTAAGATTGGCACATTCGTCACCTTCTGGA   897

Qy   620  CTTGGAACATAGTAGCTTGTTGTTCAGTGGTTCTTACTCCCAGCCTTTCTCCAAACCCTCTTAA   679
            ||||| | | ||| |||| | ||||||||| ||||   ||   | |||  |||||| | |||||
Db   898  -------------GTGGGCATTCACTCCCAAACCTGTGCTCTAGCCCTCTTAA   939

Qy   680  TTCTAGAGCAGTTGAAGACATCTCTTCACTACCATATGATAACTCAGTTATGCAATGAG   739
            |||||| ||||| ||||||| |  ||  ||| | ||| | || |||||||||| ||||||
Db   940  TTCTAGAGCAGTGGAAGACTTC---TCATCAAGGATGGATCCCTGCAGCTATGCCATGAG   996

Qy   740  TACTGAAGAAGCCAGAGATTTCTTACCTACCACATAATGTGGCCCATTAAACTTTTTTTGTCACCATC   799
            || |||||  |||||| | ||||| ||| || |||||||| ||  | |||| |||||||||||||
Db   997  TACAGAAGAGGCCCAGATTTCTTACTTCACAGTATGTGGCCTTTTAAAGTTTTTTCTGTCACCAGC   1056

Qy   800  AGAAATTGGCAAGAGCTGGTTTTTATTATATATAGGAGACCTGGGTGAGCCTGTTTGC   859
            ||||| || |||||  ||||| |||| | ||||||||| ||  |  |||| |||||||
Db  1057  AGAGCTGGCCCAGAGCTCTGATTTGCATTAGGGGCGTGGGCGTGGCGGTGGCCTGTTTTTGC   1116

Qy   860  CTGTGGGTGGGGAAGCTCAGTAACTGGGAACCAAAGGATGAGATGCTATGTCAGAACACCGGAG   919
            |||||| |||||  ||||||||  || || ||||||| | ||| | || ||||||||||||||
Db  1117  CTGTGTGTGGGGAAACTGAGCAACTGGGAACCAAAGGATGAGATGCTAGTGTCAGAGCACCGCAG   1176

Qy   920  GCATTTTCCCAACTGTCCATTTTTTGGAAAAATTCTCTAGAAAATCTGAGGTTTACATTTC   979
            |||||| || ||| |||| ||||| ||| | ||| || |||| ||| | ||||||||||||
Db  1177  ACATTTTCCCACTGTCCATTTTCTGGAAAAATACTTCAGAAACTCGCAAGTGTGCCCACAATTTC   1236

Qy   980  AAAATCTGAGCCATGCAGACACGCACATGCAGAATGAGAACATTTATGTACTGGCCATCTAG   1039
            || ||| ||| | | || | ||||||| | |||| |||||| |||  |||| | |||| |||
Db  1237  AAATCTAAGTATCCAGACACACTCGATTGAGGACATTTCTGTACTGGCCACCTAG   1296

Qy  1040  TGTTCCAGTTCACCCTGCTGACAGTGCTGGTTTTTTATTTATGTGGGGTTCGCAATGA   1099
            ||||||||||| |||| || | || |  || ||| ||| | ||  ||| ||||||||
Db  1297  TGTTCCGTTCAGCCGAGCGGTCGGAGAATCTATTACGTGGATCGGAATGA   1356

Qy  1100  TGATGTCAAAATGCTTTGTGTTGGTGATGGTGGTGTTGGGAGTGGTTGGAGTGTTGGGAGATGACCC   1159
            |||| || |||| |||| |||  || ||| || || ||| |   | || || ||||| ||||||||
Db  1357  TGATGTCAAGTGCTTTTGTTGATTGGTGAGGTGGTTGGGAACCTGACTGGAAGGGTGGAAGCTTAGTC   1416

Qy  1160  ATGGGTAGAACATCCCAAGTGGGTTTCCAAGGTGTGTGAGTTCTTGATACGAATGAAAGGCCA   1219
            | ||| |||  ||||||| | |||||||||| || || ||||| ||| | |  |||||| ||
Db  1417  CTGGATAGAACAGCCAAATGGTTTCCAAGGTGTGTGAGTTACTGGATTTGTTGTTCTT   1476

Qy  1220  AGAGTTTGTTGATGAGGAGATTCAAGGTAGGTAAGATTAGCTATGTCCTCATCTCTTCTTCTTGACTCAGCAGCTGGTTGTCAAC   1279
            |||||||||| | |||||||||||| || || |||||  | ||  | | |  || ||||||  | || || ||||||| || ||
Db  1477  GGAGTTTGTTGATGAGGATTCAAGCTAGATATCCTCATCTTCTTTGAGCAGCTGTTTGGACC   1536

Qy  1280  TTCAGATACCACTGGAGAAGAAAATGCTGACCC--------ACCAATTATTCATTTTGGACC   1333
            ||||||||||| |||| ||||||||||| |           |  ||||| |||| |||||||
Db  1537  TTCAGACACCCCAGGAGAAGAAGAAAATGCTGAGACACCTCACAGAGACAGTGGGTGCATTTTGGGCCC   1596

Qy  1334  TGGAGAAAGTTCTTCAGAAGATGCTGTTGTCATGAATACACCTGTGGTTAAATCTGCCTT   1393
            |||||| ||||  | ||| ||| || || |||| |  | |||  ||| || | |||||||
Db  1597  TGGAGAAAG---TTCGGAAGATGTCGTCATGATGATGACAGCCTT   1653

Qy  1394  GGAAATGGGCTTTTAATTAGGAGACCTGGTGAAACAAACAGTTCAAAGTAAAAATCCTGACAAC   1453
            |||| ||||| ||||| || |||| |||||||||||||||||||||| || | |||||| || |
Db  1654  GGAAATGGGCTTCAGTGACGTAGGAGCCTGGGTGAGGACAGAGGTTCAGCGCGGGCAGAATCCTGGGCAC   1713

Qy  1454  TGGAGGAGAACTATAAAACAGTTAAATGAATATGTGTCAGCACTTCTAAATGCTGAAAGATGA   1513
            ||||| ||  | ||||||||| || |||| | || ||||||||| | ||| | |  | |||||
Db  1714  TGGTGAGAACTACAGGACCGTCAATGATAATTGTCTCAGTACTTTTGAATGCTGAAGATGA   1773

Qy  1514  AAAAAGAGGAGGAGGAGAAGGAGAAAAACAAGCTGAAGAAATGGCATCAGATGATTTGTCATT   1573
            |||| ||||||||||||| ||||| |||  |||||| ||| | |||||  | || ||||  ||
Db  1774  GAGAAGGAAGAGGAGGAGGAAGAAAAGACGACTGAAGAGATGGGCATCAGCTGAGACTTATCACT   1833

Qy  1574  AATTCCGAAGAACAGAATCAGGAAATGGGCTCTCTTCTCGAACAATTGACAATGTGTTGCTTCCTATCCTGGA   1633
            |||||||||||| ||||| |||| |||||| |||| |  |||| ||||||||||| |||| |  | | ||| |
Db  1834  GATTCGGAAGAATAGAATTGGCCTTCAAAGAACTCACAGTTGACACATGCTTCCTATCCTGA   1893

Qy  1634  TAATCTTTTAAAGGCCAATGTAATTGAATTCCAACACAGAAGAACATGATATATATTATAAAACAAAAAC   1693
            ||||  ||||| |||| ||||| || ||||||||| | ||||| | |  | | |||| |  ||| ||||
Db  1894  TAAATCTTCTTGAGGCCCAGTGTAATTACAAACACATGGATTTCAGGCTTGTCATTGCTTAGCAGAAAAC   1953

Qy  1694  ACAGATACCTTTACAGCGAGGAGAACATTGATATCCATTTTGGGTAAAGGGAAAATGCTGC   1753
            |||||| |||| ||| |||||| |||| | |||| ||||||||  |||||||| || ||||
Db  1954  ACAGATACCCTTACAGCAAAGAGAGCTTATTGACACCGTTTTAGTCAAGGGAAAATGCTGC   2013

Qy  1754  GGCCAACATCTTCAAAAAACTGTCTAAAAGAAATTGACTCACATTGTAATAAGAAACTTATT   1813
            |||||||||||||| || |  | |  | |||||||||||||| | ||||| ||||| |||||
Db  2014  AGCCAAACATCTTCAAAAAACTCTCTGAACATTGACTCCACGTTATAGGTCCTATCCTCGA   2073

Qy  1814  TGTTGGATAAGAATATGAAGTGTAATTCCAACACAGAAGAATGTTTCAGGTCTGTCACTGGAGA   1873
            || || |||| | | ||  ||| || || ||| | ||||| | || ||||| |||| ||||
Db  2074  TGTGGAAAAGGAATATGAAGTGTAATTCCAACACATGATGATGGTTCAGGTGTTCATTGGAGGA   2133

Qy  1874  ACAATGAGGAGGTTGCAAGAACAAGCAGAACTGTAAAAGTGGTGTATGGGACAAAAGAAGTTTC   1933
            |||| |||||||  ||||||| | |||||||||| | ||||| | ||||| ||||||| ||||
Db  2134  GCAGTTGCGGAGAATTACAAGACCAACACTTGCAAAAGTGTAATGGACAGAGGAGGGTTTC   2193

Qy  1934  TGTTGTATTTTATTCCTTGGTCATCTGGTATGCCCAGGAAGTGTTGCCCCTTCTCTAAG   1993
            |||| | ||||| |||| ||  | | ||| || |||||| | |||| ||||| |||||
Db  2194  TATTGTTGTCATTCCGTGTGGTCATCTACTAGTCTGCCAGAAATGTGCCCTTCTCTAAG   2253

Qy  1994  AAAATGCCCTATTTGCAGGGGGTATAACAAGGGTACTTGGTTCGTACATTTCTCTCTTAAAG   2053
            || || ||||| || |||||| | | || |||| | || || |  | ||||||| |||||
Db  2254  GAAGTGCCCATCTGCAGGGGCAGCAATCAAGGGACTGTGCCGCACATTTCTTCTCCATGA--   2311

Qy  2054  AAAAATAGTCTATATTTTTTAACCTGCATAAAAAGGTCTTTAAAATATTGTTGAACACTTGA   2113
            |||||| ||| | | |  |||||| ||| ||||||| | || || |||| |||||  |||||
Db  2314  --------------GTGAAGAATGGTCTGAAAGTATTGTTGGACATACAGA   2347

Qy  2114  AGCCATCTAAAGTAAAAGGGAATTATGAGTTTTTCAAATAACATTCATGTTTCTTAGT   2173
            || || || | ||||| || | | ||   ||| | |||| |||| || | |||  |
Db  2348  AGCTGTGCAGAACAAAAGAATGAACTACTGA---TTTCAGCTCTTCAGGCAGGACATTTCACT   2404

Qy  2174  CTGCTTTGGTACTAATAATATCTTGTTTTCTGAAAAGATGGGTATCATATATTTTAATCTTAATC   2233
            ||| |  ||| |||| | | | || ||| |  ||||||| ||| || ||||| | |||| ||
Db  2405  CTCTTTCAAGATTAGTTAATCTTGCTTTAATGAAAGGGTAGCATTGTAGTTATATATTTTAAGCTTAGTC   2464

Qy  2234  TGTTTATTTACAAGGGAAGAGATTTAATGTTTGGGTGAACTATTAGTAATGGTAATGTGTACCTA   2293
            ||  | ||| ||||| ||||| | | ||  || || |||| | || | |  | || ||| ||||
Db  2465  TG-----TTGCAAGGGAAGGGTCTATG--CTGTTGAGGCTACAGGACTGTGCTCGTTGTTCCAGAG   2518

Qy  2294  AGGGAGTAGTGTGTCACTCGCTTGTGTTATGCATCATCATTCAGGAGTTACTGGATTTGTTGTTCTT   2353
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                  ||||| | |   ||   |||    |     ||||||| ||  |||||   |||   ||
Db         2519 CAGGAGTTGGGATGCTTGCTGTATGTC----CTTCAGGACTTCTTGGAT---TTGGAATT 2571

Qy         2354 TCAGAAAGCTTTGAATACTAAATTATAGTGTAGAAAAGAACTGGAAACCAGGAACTCTGG 2413
                |  |||||||||| || |         |      | | | |  ||||||||   ||||||
Db         2572 TGTGAAAGCTTTGGATTCAGGTGATGTGGAGCTCAGAAATCCTGAAACCAGTGGCTCTGG 2631

Qy         2414 AGTTCATCAGAGTTATGGTGCCGAATTGTCTTTGGTGCTTTTCACTTGTGTTTTAAAATA 2473
                   |||    ||||| ||| ||   |  |  ||||||||||||  || ||    ||||||
Db         2632 TACTCA--GTAGTTAGGGTACCCTGTGCTTCTTGGTGCTTTTCCTTTCTG---GAAAATA 2686

Qy         2474 AGGATTTTTCTCTTATTTCTCCCCCTAGTTTGTGAGAAACATCTCAATAAAGTGCTTTAA 2533
                |||||||||||  || |  |     |  | |||  |  | |  |  || ||    |||
Db         2687 AGGATTTTTCTGCTACTGGTAAATATTTTCTGTTTGTGAGAAATATATTAAAGTGTTTCT 2746

Qy         2534 AAAAAA 2539
                   |||
Db         2747 TTTAAA 2752

RESULT 11
US-09-974-592-13
; Sequence 13, Application US/09974592
; Patent No. US20020120121A1
; GENERAL INFORMATION:
;  APPLICANT: Korneluk, Robert G
;  APPLICANT:  MacKenzie, Alexander E
;  APPLICANT:  Liston, Peter
;  APPLICANT:  Baird, Stephen
;  APPLICANT:  Tsang, Benjamin K
;  APPLICANT:  Pratt, Christine
;  TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
;  TITLE OF INVENTION:  NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
;  TITLE OF INVENTION:  DISEASE
;  FILE REFERENCE: 07891/009004
;  CURRENT APPLICATION NUMBER: US/09/974,592
;  CURRENT FILING DATE:  2001-10-09
;  PRIOR APPLICATION NUMBER: US 09/617,053
;  PRIOR FILING DATE: 2000-07-14
;  PRIOR APPLICATION NUMBER: US 08/800,929
;  PRIOR FILING DATE: 1997-02-13
;  NUMBER OF SEQ ID NOS: 17
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 13
;   LENGTH: 3151
;   TYPE: DNA
;   ORGANISM: Mus musculus
US-09-974-592-13

  Query Match             51.9%;  Score 1343.6;  DB 10;  Length 3151;
  Best Local Similarity   76.2%;  Pred. No. 6.7e-273;
  Matches 1852;  Conservative    0;  Mismatches  499;  Indels   79;  Gaps   13;

Qy          116 TTTGGTGGTAAAAATCTTAGTTCATGTGAAGAAATTTCATGTGAATGTTTTAGCTATCAA 175
                | ||||||  |     |       ||   ||||| ||||| || | ||||  |||  |||
Db          702 TGTGGTGGAGATCTAGTGTCCAAGTGGTGAGAAACTTCATCTGGAAGTTTAAGCGGTCAG 761

Qy          176 ACAGCACTGTCACCTACTCATGCACAAAACTGCCTCCCAAAGACTTTTCCCAGGTCCCTC 235
                | |      |   ||||||||| ||||||||| |||||| |||||   || |||| |||
Db          762 AAATACTATT--ACTACTCATGGACAAAACTGTCTCCCAGAGACTCGGCCAAGGTACCTT 819

Qy          236 GTATCAAAACATTAAGAGTATAATGGAAGATAGCACGATCTTGTCAGATTGGACAAACAG 295
                  | |||||  ||||  ||||||||||  | ||||| |||||||| ||||||||||
Db          820 ACACCAAAAACTTAAACGTATAATGGAGAAGAGCACAATCTTGTCAAATTGGACAAAGGA 879

Qy          296 CAACAAACAAAAAATGAAGTATGACTTTTCCTGTGAACTCTACAGAATGTCTACATATTC 355
                 | | || |||||||||||| ||||||||| ||||||||||||| ||||||||||||||||
Db          880 GAGCGAAGAAAAAATGAAGTTTGACTTTTCGTGTGAACTCTACCGAATGTCTACATATTC 939

Qy          356 AACTTTCCCCGCCGGGGTGCCTGTCTCAGAAAGGAGTCTTGCTCGTGCTGGTTTTTATTA 415
                | |||| |||   || || ||||||||||| |||||||| ||||||||||| ||||||||
Db          940 AGCTTTTCCCAGGGGAGTTCCTGTCTCAGAGAGGAGTCTGGCTCGTGCTGGCTTTTATTA 999

Qy          416 TACTGGTGTGAATGACAAGGTCAAATGCTTCTGTTGTGGCCTGATGCTGGATAACTGGAA 475
                ||| |||||||||||||| ||||| ||||||||| |||||||||||| |||||||||||||
Db         1000 TACAGGTGTGAATGACAAAGTCAAGTGCTTCTGCTGTGGCCTGATGTTGGATAACTGGAA 1059

Qy          476 ACTAGGAGACAGTCCTATTCAAAAGCATAAACAGCTATATCCTAGCTGTAGCTTTATTCA 535
                || ||| ||||||||| || ||||||| | |||| | |||| ||||| ||||| | ||
Db         1060 ACAAGGGGACAGTCCTGTTGAAAAGCACAGACAGTTCTATCCCAGCTGCAGCTTTGTACA 1119

Qy          536 GAATCTGGTTTCAGCTAGTCTGGGATCCACCTCTAAGAATACGTCTCCAATGAGAAACAG 595
                || |||| |||||| ||||||   ||  | ||||||||||| ||||||  ||| ||  ||
Db         1120 GACTCTGCTTTCAGCCAGTCTGCAGTCTCCATCTAAGAATATGTCTCCTGTGAAAAGTAG 1179

Qy          596 TTTTGCACATTCATTATCTCCCACCTTGGAACATAGTAGCTTGTTCAGTGGTTCTTACTC 655
                 |||||||||||| | |||        |                   ||||   | ||||
Db         1180 ATTTGCACATTCGTCACCTCTGGAACGAG------------------GTGGCATTCACTC 1221

Qy          656 CAGCCTTTCTCCAAACCCTCTTAATTCTAGAGCAGTTGAAGACATCTCTTCATCGAGGAC 715
                || ||| |   | | ||||||||||||||||||||| |||||| ||   ||||| ||||
Db         1222 CAACCTGTGCTCTAGCCCTCTTAATTCTAGAGCAGTGGAAGACTTC---TCATCAAGGAT 1278

Qy          716 TAACCCCTACAGTTATGCAATGAGTACTGAAGAAGCCAGATTTCTTACCTACCATATGTG 775
                  | |||| ||| ||||| ||||||| ||||| |||||||||||||| |||  ||||||
Db         1279 GGATCCCTGCAGCTATGCCATGAGTACAGAAGAGGCCAGATTTCTTACTTACAGTATGTG 1338

Qy          776 GCCATTAACTTTTTTGTCACCATCAGAATTGGCAAGAGCTGGTTTTTATTATATAGGACC 835
                ||| |||| |||| |||||||| ||||  |||| |||||||| || ||||| ||||| ||
Db         1339 GCCTTTAAGTTTTCTGTCACCAGCAGAGCTGGCCAGAGCTGGCTTCTATTACATAGGGCC 1398

Qy          836 TGGAGATAGGGTAGCCTGCTTTGCCTGTGGTGGGAAGCTCAGTAACTGGGAACCAAAGGA 895
                |||||| ||||| ||||| ||||||||||||||||| ||  || ||||||||||||||||
Db         1399 TGGAGACAGGGTGGCCTGTTTTGCCTGTGGTGGGAAACTGAGCAACTGGGAACCAAAGGA 1458

Qy          896 TGATGCTATGTCAGAACACCGGAGGCATTTTCCCAACTGTCCATTTTTGGAAAATTCTCT 955
                ||||||||||||||| |||| | || ||||||||| ||||||||||| |||||||| ||
Db         1459 TGATGCTATGTCAGAGCACCGCAGACATTTTCCCCACTGTCCATTTCTGGAAAATACTTC 1518

Qy          956 AGAAACTCTGAGGTTTAGCATTTCAAATCTGAGCATGCAGACACATGCAGCTCGAATGAG 1015
                |||||| | ||||||||| || ||||||||| || |||||||||||  | ||||| ||||
Db         1519 AGAAACACAGAGGTTTAGTATATCAAATCTAAGTATGCAGACACACTCTGCTCGATTGAG 1578

Qy         1016 AACATTTATGTACTGGCCATCTAGTGTTCCAGTTCAGCCTGAGCAGCTTGCAAGTGCTGG 1075
                 |||||| ||||||||||| ||||||||||| |||||||| ||||||||||||||||||||
Db         1579 GACATTTCTGTACTGGCCACCTAGTGTTCCTGTTCAGCCCGAGCAGCTTGCAAGTGCTGG 1638

Qy         1076 TTTTTATTATGTGGGTCGCAATGATGATGTCAAATGCTTTTGTTGTGATGGTGGCTTGAG 1135
                 || ||||| |||| ||||||||||||||||||| ||||||||||||||||||||||||||
Db         1639 ATTCTATTACGTGGATCGCAATGATGATGTCAAGTGCTTTTGTTGTGATGGTGGCTTGAG 1698

Qy         1136 GTGTTGGGAATCTGGAGATGATCCATGGGTAGAACATGCCAAGTGGTTTCCAAGGTGTGA 1195
                 ||||||||| ||||||||| || ||| ||||||| |||||| |||||||||||||||||
Db         1699 ATGTTGGGAACCTGGAGATGACCCCTGGATAGAACACGCCAAATGGTTTCCAAGGTGTGA 1758

Qy         1196 GTTCTTGATACGAATGAAAGGCCAAGAGTTTGTTGATGAGATTCAAGGTAGATATCCTCA 1255
                ||||||||||||| ||||| || || |||||||||||||||||||||||| ||||||||||||
Db         1759 GTTCTTGATACGGATGAAGGGTCAGGAGTTTGTTGATGAGATTCAAGCTAGATATCCTCA 1818

Qy         1256 TCTTCTTGAACAGCTGTTGTCAACTTCAGATACCACTGGAGAAGAAAATGCTGACCC--- 1312
                ||||||||| ||||||||||| |||||||| ||| | ||||||||||||||||||||
Db         1819 TCTTCTTGAGCAGCTGTTGTCCACTTCAGACACCCCAGGAGAAGAAAATGCTGACCCTAC 1878

Qy         1313 ---ACCAATTATTCATTTTGGACCTGGAGAAAGTTCTTCAGAAGATGCTGTCATGATGAA 1369
                     || |  | ||||||||| ||||||||||||   |||  ||||||  |||||||||
Db         1879 AGAGACAGTGGTGCATTTTGGCCCTGGAGAAAG---TTCGAAAGATGTCGTCATGATGAG 1935

Qy         1370 TACACCTGTGGTTAAATCTGCCTTGGAAATGGGCTTTAATAGAGACCTGGTGAAACAAAC 1429
                 || |||||||||||| | ||||||||||||||||| | |||   ||||||||| ||| ||
Db         1936 CACGCCTGTGGTTAAAGCAGCCTTGGAAATGGGCTTCAGTAGGAGCCTGGTGAGACAGAC 1995

Qy         1430 AGTTCAAAGTAAAATCCTGACAACTGGAGAGAACTATAAAACAGTTAATGATATTGTGTC 1489
                 |||||  |  | |||||| | ||||| |||||||| |  || || ||||||||||| ||
Db         1996 GGTTCAGCGGCAGATCCTGGCCACTGGTGAGAACTACAGGACCGTCAATGATATTGTCTC 2055

Qy         1490 AGCACTTCTAAATGCTGAAGATGAAAAAAGAGAGGAGGAGAAGGAAAAACAAGCTGAAGA 1549
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RESULT 12

US-09-201-936-41
; Sequence 41, Application US/09201936
; Publication No. US20020187946A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: MacKenzie, Alexander E.
; APPLICANT: Baird, Stephen
; APPLICANT: Liston, Peter
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
; TITLE OF INVENTION: PROBES, AND DETECTION METHODS
; FILE REFERENCE: 07891/003003
; CURRENT APPLICATION NUMBER: US/09/201,936
; CURRENT FILING DATE: 1998-12-01
; EARLIER APPLICATION NUMBER: 09/011,356
; EARLIER FILING DATE: 1998-02-04
; EARLIER APPLICATION NUMBER: PCT/IB96/01022
; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 08/576,956
; EARLIER FILING DATE: 1995-12-22
; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 41
; LENGTH: 2416
; TYPE: DNA
; ORGANISM: Mus musculus
US-09-201-936-41

Query Match   51.4%;   Score 1330.4;   DB 10;   Length 2416;
Best Local Similarity   76.4%;   Pred. No. 3.5e-270;
Matches 1816; Conservative   0; Mismatches 486; Indels 74; Gaps 12;

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Qy  116  TTTGGTGGTAAAAATCTTAGTTCATGTCATGTGAAGAAATTTCATGTGAATGTTTTAGCTATCAA  175
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Db    2  TGTCGTGGGAGATCTATTGTCCAAGTGGTGAGAAACTTCATCTGGAAGTTTAAGCGGTCAG       61

Qy  176  ACAGCACTGTCACCTACTCATCACAAAAACTGCCTCCCAAAGACTTTTCCAGGTCCCTC        235
         || ||||     |||||||||| ||||||||||||||||||||||||||||||||||||
Db   62  AAAATACTATT--ACTACTCATGGACAAAAACTGTCTCCCAGAGACTTACCTTT              119

Qy  236  GTATCAAAACA-TTAAGAGTATAATGGAAGATAGCACGATCTGTCAGATAGGACAAAACA        294
         |||||||||| ||||| ||||||||||||||||| |||| |||||||||||||||||
Db  120  ACACCCAAAAACTTAAACGTATAAATGGAAGAGACACAATCTTGTCAAATTGGACAAAGG        179

Qy  295  GCAACAAACAAAAAAATGAAGTATGACATTTTCCGTGTGAACTCTACAGAAGTCTACACATATT  354
         ||||| ||||||||||||||||| |||||||| |||||||||||| |||| ||||||||||||
Db  180  AGAGCGAAGAAAAAATGAAGTTTGACATTTTCGTGTGTGAACTCACCGAATGTCTACACATATT  239

Qy  355  CAACTTTTCCCGCCGGGGGTGCCTGTCTCAGAAAAGGAGTCTTGCCTGTCTCGGTTTTTTATT  414
         || |||||||| ||||| ||||||||||||||||||||||||||||||| ||||||||||||
Db  240  CAGCTTTTCCCAGGGGAGTTCCTGTCTCAGAGAGGAGTCTTGGCTGCTCGGCTGTTTTATT     299

Qy  415  ATACTGGTTGTGAAATGACAAGGTCAAAATCCTTCTCGTTGTGGCCTGATGTTGGATAAACTGGA  474
         |||| ||||||||||||||||| |||| |||||||||||||||||||||||||||||||||||
Db  300  ATACAGGGTGTGAAATGACAAAGTCAAGTGCTTCGGCTGTGGCCTGATGTTGGATAAACTGGA     359

Qy  475  AACTAGGAGGACAGTCCTATTCAAAAGCATAAACAGCTATATCCTAGCTGTAGCTTTATTC  534
         || |||| ||||| |||| ||||||| ||||||||
Db  360  AACAAGGGGGACAGTCCTTGTTGAAAAGCACAGACAGTTCTATCCCAGCTGCCAGCTTTGTAC  419

Qy  535  AGAATCTGGGTTTCAGTGACCACCTCTAAGAATACGTCTCCAATGAGAAACA              594
         |||| |||||| |||
Db  420  AGACTCTGCTTTCAGCCAGTCTCGCAGTCCATCTAAGAATAATAGTCTCCTGTGAAAAGTA  479

Qy  595  GTTTTGCACATTCATTATCTCCCACCTTGGAACATAGTAGCTTGTTGTTCAGTGGTTCTTACT  654
         ||||||| ||||      |||
Db  480  GATTTGCACAATTCGTCACCTCTCGGAACGAG                                 521

Qy  655  CCAGCCTTTTCTCCAAACCCTCTTAATTCTTAGAGCAGTTGAAGACATCTTCATCGAGGA  714
         ||| ||||   |||
Db  522  CCAACCTGTGCTCTAGCCCTTCTTAATTCTTAATTCTTTAGAGCAGTGGAAGACTTC---TCATCAAGGA  578
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Qy  2056  AGTACTTTTGAATGCTGAAGATGAGGAGAGAGAGAGGAGGAGGAGGAGGAGAAAGACAGAGA       2115
          |||||      ||||  |||
Db  1550  AATGGCATCAGATGATTTGTCATTAATTCGGAAGACAGAATGAGGCTCTCTTCTTTCAACATT   1609

Qy  2116  GACACGTGTGCTTCCTATCCTCGGATAATCTTTAAAGGCCAATGTAATTAATAAACAGGA       2175
          |||||
Db  2176  GACACATGTCCTTCCTATCCTCGGATAAATCTTCTTGAGGCCAGTGTGTAATTACAAAACAGGA  2235
```

Qy 715 CTPAACCCCTACAGTTATGCAAATGACTGAAGAAGCCAGAGATTTCTTACCTACTACCATATGT 774
        ||||| |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 579 TGGATCCCTGCAGCTTATCAGTCGCATAGCCATGAGTACAGAAGAGGCCAGATTTCTTACTTACAGTATGT 638

Qy 775 GGCCAATPAACTTTTTTTGTCACCCATCAGAATTGGCACAGAAAGACTGGTTTTTTTATTATATAGGAC 834
        |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 639 GGCCTTTAAGTTTTCTTGTCACCAGCAGAGACTGGTTTTTTTATTATATAGGGC 698

Qy 835 CTGGAGATAGGGTAGCCTGCTTGCCCTGTGGGGGAAGCTCAGTAACTGGGGAACCAAAAGG 894
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db 699 CTGGAGACAGGGGTGGCCCTGTTTTGCCTGTGGGGGGAAACTGACGAACCCAAAAGG 758

Qy 955 TAGAAACTTCTGAGGTTTAGCATTTCAAAATCTGAAATCTGAGCATGCAGACACATGCAGAGCTGAAATGA 1014
        ||||||||||||||||||||||||||||||||||||||||||||
Db 819 CAGAAAACACAGAGGTTTAGTTATATGTATATCAAAAATCTPAAGTATCAGAACACACTCTGATTGA 878

RESULT 13
US-09-954-456-1635
Sequence 1635, Application US/09954456
Patent No. US20020115057A1
GENERAL INFORMATION:
APPLICANT: Young, Paul
TITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using Cance
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-76
CURRENT APPLICATION NUMBER: US/09/954,456
CURRENT FILING DATE: 2001-09-18
PRIOR APPLICATION NUMBER: US/60/233,617
PRIOR FILING DATE: 2000-09-18
PRIOR APPLICATION NUMBER: US/60/234,052
PRIOR FILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: US/60/234,923
PRIOR FILING DATE: 2000-09-25
PRIOR APPLICATION NUMBER: US/60/235,134
PRIOR FILING DATE: 2000-09-25
PRIOR APPLICATION NUMBER: US/60/235,637
PRIOR FILING DATE: 2000-09-26
PRIOR APPLICATION NUMBER: US/60/235,638
PRIOR FILING DATE: 2000-09-26
PRIOR APPLICATION NUMBER: US/60/235,711
PRIOR FILING DATE: 2000-09-27

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; PRIOR APPLICATION NUMBER: US/60/235,720
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: US/60/235,840
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: US/60/235,863
; PRIOR FILING DATE: 2000-09-27
; NUMBER OF SEQ ID NOS: 2276
; SOFTWARE: Patentin version 3.0
;
; SEQ ID NO 1635
; LENGTH: 3076
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-954-456-1635

Query Match   44.7%;   Score 1157;   DB 10;   Length 3076;
Best Local Similarity   77.3%;   Pred. No. 1.4e-233;
Matches 1474;   Conservative   0;   Mismatches 415;   Indels 18;   Gaps 5;
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Qy    170 TATCAAACAGCACTGTCACCTACTCATGCACAAAACTGCCTCCC--------AAAGACTTTT 223
          |||| ||| ||  ||| ||||||||||||| || ||||| |||       |||||||||
Db    640 TATTAAAACTGATAAAAGCAAGCCATGCACAAAACTACCTCCCTAGACAAAAGGCTAGTC 699

Qy    224 CCCAGGTGCCCTCGTATCAAAAACATTAAGAGTATATAATGGACAAGAGCACGATCCTGTGTCAGA 283
          ||||| |||||||| ||| |||||| ||  ||| || || ||| || ||||  ||||| ||||||
Db    700 CCTTTTCTTCCCCATTCATTTCATGTAGAACATAGTAGAAAACAGCATATTCTTATCAAA 759

Qy    284 TTGGACAAAACAG---CAACAAACAAAAATGAAGTAATGAGCTTGTCAGTAACTTCTTACAG 340
          |||||||| ||    |||| ||| || ||| || |||  ||||||  ||| |||| ||||
Db    760 TTTGATGAAAAGGGCCACACAGTTTGAAATACGGACTTTGTCATGTGAACTGTACCG 819

Qy    341 AATGTCTACATATTCAACTTTTCCCCGCCCCGGGGTGCCTGTCTCAGAAAGGAGTCTTGCTCG 400
          |||| ||  |||| ||||||  || |||  || |  || || ||||||||||| |||| ||
Db    820 AAATGTCTACGTATTCCACTTTTCCTGCTGGGGGGTTCCTGTGTCTCAGAAAAGGAGTCTTGGCTCG 879

Qy    401 TGCTCGGTTTTTATTATTATACTGGTGTGAATGACAAGGTCAAAATGCTTCTGTTGTTGTTGGCCTGAT 460
          ||||||| || || ||||| ||  ||||||| ||||| || || ||||||| || |||||||  |||
Db    880 TGCTCGGTTTCTATTACACTGGTGTGCAAGGTCAAAATGCTTCTGTTGTTGTGGCCTGAT 939

Qy    461 GCTGGATAACTGAAAACTAGGACAGTCCTTCTTATTCTCAGAAAGCATAAAACAGCTATATCCTAG 520
          |||||| |||||| |||||  |  ||||| ||| ||| ||||| ||| || |||| || ||| ||
Db    940 GCTGGATAACTGGAAAAGAGGAGACAGTCCTTACTGAAAAGCATAAAAAGTTGTATCCCTAG 999

Qy    521 CTGTAGCTTTATTCAGAATCTGGTTTCAGCT---AGTCTGGGATCCACCTTCTTAAGAAATAC 577
          |||||||| ||  ||||| || ||| ||||    |||| |||||||| |||| || |||||||
Db   1000 CTGCAGAATTCGTTCAGAGGTCTAAAATTCCGTTAACAACTTGGAAGCTACCTCTCAGCCTAC 1059

Qy    578 GT---CTCCAATGAGAAACAGTTTTGCACATTCATTTATCTCCCACCTTGGGACACATAGTAG 634
          ||   ||||| || ||||||| | |||||| ||||||| || || |||||||||| ||| ||
Db   1060 TTTTCCTTCTTCAGTAACAAAATTCCACACACTCATTACTTCCGGGTACAGAAAACAGTGG 1119

Qy    635 CTTGTTCAGTGGGTTCTTACTCCAGCCTTTCTCCAAACCCTCTTAATTCTTAGAGCAGTTGA 694
          ||||||| ||||| |||| ||||||||| ||| ||| || ||  || || ||| ||||| |
Db   1120 ATATTTCCGGTGGGCTCTTATTCAAAACTCTCCATCAAATCCGTAAACTGGAGAGCAAAATCA 1179

Qy    695 AGACATCTCTTCATCGGAGGACTAACCCCCTACAGTTTATTGCAATGGAAGAAGGCCAG 754
          ||||| |||| ||| |||||| |||||||| ||||   || ||| | |||||||| |||
Db   1180 AGATTTTTTCTGCCTTGATGAAGTTCCTACCACTGTGCAATGACTGGGTGGGTGGGAGC 1239

Qy    755 ATTTCTTACCTACCATATGTGGCCATTAACTTTTTTGTCACCATCAGAAATGGCAAGAGC 814
          ||||| |||  ||||||| ||| ||||| ||||   | ||| ||||||| ||||| |||
Db   1240 ATTACTTACTTTTCAGACATGGCCATTGACTTTTCTGTCGCAAACAGATCTGGCAAAAGC 1299

Qy    815 TGGGTTTTTATTATATAGGACCTGGAGATAGGGTAGCCTGCTTTGCCTGTGGGGGGAGCT 874
          ||||| ||| ||| ||  ||||| ||| |||||| |  || ||| ||||||| || |||
Db   1300 AGGCCTTTTTACTACAGACGAGCACAGACGTGGCCTTG|TTTTGCCTGTGTGGTGGGAAATT 1359

Qy    875 CAGTAACTGCGGCGAACCAAAAGGATGATGCTATGTCACGGAGGACCATTTTCCCAACTG 934
          |||||||   |||  | || || ||| ||||| || | ||| || ||||||| ||||
Db   1360 GAGCAATTGGGGACCGGAAGGGATAAGTCTAAGTCAAGACACACTGAGACCATTTTCCCAAATG 1419

Qy    935 TCCATTTTTTGGAAAAT---TCTCTAGAGACACTTCAAAATCTGAGCAT 991
          ||| ||||| ||||| |   |||| |||||||| || || || |||||
Db   1420 CCCATTTTATAGRAAAATCAGCTTCAAGACACACTGACGATACACAGTTTCTTAATCTGAGCAT 1479
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Qy    992 GCAGACACAATGCAGCTCGAAATGAGAAACATTTATGTGTTTACTGTTTCCAGTTCA 1051
          ||||||| ||||||||||||||| |||| ||||||| ||| ||| |||| ||||||
Db   1480 GCAGACACATGCAGCCCCGCTTTAAAAACATTCTTTAACTGGCCCTTCAGTTAA 1539

Qy   1052 GCCTGAGCAGCTTGCAAGTGCTGGTTTTTTTATTATGTGGTGTCGCAAATGATGAGTGTCAAATG 1111
          |||||||||||||||||||| ||||   |||||| ||||||||||| ||||| || ||||||||
Db   1540 TCCTGAGCGGCTTGCAAGTGCCGGGGTTTTTTTATTAGTGGGGTTAACAGTGATGAGTCAAAATG 1599

Qy   1112 CTTTTTGTTGTGATGGGTGGCGTTGAGGTGTTGGGAATCTGGAGTGATCCATGGGTAGAACA 1171
          |||||||||| |||||||| | || |||||||||||||||| || || |||  ||||| ||
Db   1600 CTTTTTGCTGTGATGGGTGACTCAGGTGTTGGGAAATCTGAGATGATCCATGGGTTCAAACA 1659

Qy   1172 TGCCAAGTGGTTGTTTCCAAGGTGTGAGTTCTTGATGTAGGAATGAAATGGTTAGTTGTTGA 1231
          ||||||||||||||  ||||||||| ||||||||||||| ||||| || ||||  |||||||
Db   1660 TGCCAAGTGGTTTTCCAAGTGTGAGTACTGACGTAAGAAATTAAAGGACCAGGAGTTCATCCG 1719

Qy   1232 TGAGATTCAAGGTAGAATATCCTCCATCTTCCTTGAAACAGCCTGTTGTCGCAACTTCAGATACCAC 1291
          |||| ||||||| ||| ||| ||||||| ||  ||||||||  |||||||| |||||| ||||||
Db   1720 TCAAGTTCAAGCCCAGTTACCCTCCATCTCACTTGAACAGCTGTCTCCACATCCAGCAGCCC 1779

Qy   1292 TGGAGAAGAAAAATGCTGACCCACCAATTATTCATTTTGGACCTGGAGAAAGTTCTTCAGA 1351
          |||||||||||| ||| |||| || || ||||||||| |||| || ||||||  ||||||
Db   1780 AGGGAGATGAAAAATGCAGAGAGTCATCAAATTATCCATTTTGAACCTGGAGAAGACCATTCAGA 1839

Qy   1352 AGATGCTGTCATGATGAATGAATACACCCTGTGTTAAATCTGCCTTGGAAATGGCGCTTTAATAG 1411
          ||||||| |||||||||| || || || || |||  | ||||| ||||| |||| |  || |||
Db   1840 AGATGCAATCAGATGAATACTCCTGTGATTAATGCTGGAAATGGGCTTTAGTAG 1899

Qy   1412 AGACCTGGTGAAACAAAACAGTTCAAAGTAAAAATCCTGACAACCTGGAGGAGAACTATAAAAC 1471
          |||| ||| |||| ||  |||||  ||||| |||| |||| ||| ||| || ||||| ||||
Db   1900 AAGCCTGGTAAAAACAGGACAGTTCAAAAGAAAAACCTAGCAACTGGGAGGAATATTAGAAT 1959

Qy   1472 AGTTAATGATAATTGTGTCAGCACACTTCTAAATGCTGAAAGATGAAAAAAGAGGAGGAGGAGAA 1531
          ||||||||| || |||| ||||| |||  || || |||  ||| || ||  |||| || || ||
Db   1960 AGTCAATGATCTTTGCTTTGACTTTACCATGTGTAAATTAAAGGATGAAGATGAAGGAGAGG 2019

Qy   1532 GGAAAAACAAGCTGAAGAAAATGGCATCAGATGAGATTTGTCATTAATTCGGAAGGAACAGAAT 1591
          |||||| ||||| | |||||| || |||| || |||| |||  ||||| ||||  |||| |||
Db   2020 AGAAAGAGCAACTGAGGAAAAAAGAAATCAAAATGATTATTATTAAATCCGGAAGAAATAGAAT 2079

Qy   1592 GGCTCTTCTTCAACAATTGACACATGTGGTCGCTCCTCCTGGATAATCTTTTAAAGGCCAA 1651
          |||| |||| ||| |  ||| || |||| |  || || |||| || |||| | ||||| ||
Db   2080 GGCACTTTTTCAACATTTGACATTTGGTAATTCCAATCCTGGATAGTCTTACTGCCCG 2139

Qy   1652 TGTAATTAATAAAACAGGAACAATGCATCAGATGATTTGTCATTAATTGTAGAACAGAAGC 1711
          ||  ||||| || || || ||| ||| | |||| | ||||  |||| ||||  ||||| ||
Db   2140 AATTAATTAATGAACAACAGAAACATGAATGTTATTTAAACAGAAATGTCTCTTTACAAGC 2199

Qy   1712 GAGAGAACTGATTGATACCATTTTGGGTTAAAGGAAAAATGCTGGCGGCCAACATCTTCAAAAA 1771
          |||| |||| |||| |||| ||||||||| |||||| ||| ||| ||| ||||| ||||| |||
Db   2200 AAGAGAACTGAATTGAATAGCATGTTAGTAGGAGAAATATTGCAGCCACTGTAATCAGAAAA 2259

Qy   1772 CTGTGTCTAAAAGAAAATTGACTCTACACATTGTATAAGAACTTATTTGTGGATAAGAATATGA 1831
          || | || ||||| || || || || ||| | |||| ||| | |||||||| |||| |||||
Db   2260 CTCTCTGCAAGAAGCTGAAGCTGAAGCCCACTTTAATTGTCAAACAGGACATAA 2319

Qy   1832 GTATATTCCAACAGAAGAGATGTTTCAGGTCTCACTGCGGAAGAACAAATTGAGGAGGTTGCA 1891
          |||| ||| |||||||| || |||||||| || |||||| || || || |||| ||  ||||
Db   2320 ATATATTCCCACAGAAGAGTTTCAGACTACCAGTGGAAAACAAATTGCGCAGGGACTACA 2379

Qy   1892 AGAAGAAACTTGTAAAGTGTGTATGGACAAAGAAGAGTTTCGTGTTGTATTTATTCCTTG 1951
          |||||||| || ||||| ||| || || ||||||||  |||||| ||  || ||||| ||
Db   2380 AGAAGAAAGAACATGTTAAAGTGTATGGACAAAGAACAATTGCCACTGTTTTTATTCCTTG 2439

Qy   1952 TGGTCATCTGGTAATGCCAGGAAATGTGCCCCTTCTCTAAGAAAAATGCCCTATTTTGCAG 2011
          ||||||||||| |  ||| ||| |||  | || |||  |||||||| |  |||||| |||
Db   2440 TGGTCATCTAGTAATGCAAAGATTCTGTGCTCTTCTTTTAAGAAAGTGTCCTATTTGTAG 2499

Qy   2012 GGGTATAATCAAAGGGTACTGTTGCTACAGTTCGGTACATTTCTTCTTAAGAAAAAA 2058
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Db   2500 GAGTACAAATCAAAGGGTACAGTTCGGTACACATTTCTTTCATGGAAGAGA 2546
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RESULT 14
US-09-954-531-16
; Sequence 16, Application US/09954531
; Patent No. US2002016518OA1
; GENERAL INFORMATION:
; APPLICANT: Weaver, Zoe
; TITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using Canc
; TITLE OF INVENTION: Gene Sets
; FILE REFERENCE: 689290-77
; CURRENT APPLICATION NUMBER: US/09/954,531
; CURRENT FILING DATE: 2002-05-02
; PRIOR APPLICATION NUMBER: US/60/233,133
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: US/60/234,009
; PRIOR FILING DATE: 2000-09-20
; PRIOR APPLICATION NUMBER: US/60/234,034
; PRIOR FILING DATE: 2000-09-20
; PRIOR APPLICATION NUMBER: US/60/234,509
; PRIOR FILING DATE: 2000-09-22
; PRIOR APPLICATION NUMBER: US/60/234,567
; PRIOR FILING DATE: 2000-09-22
; NUMBER OF SEQ ID NOS: 1392
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 16
; LENGTH: 3076
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-954-531-16

Query Match   44.7%; Score 1157; DB 10; Length 3076;
Best Local Similarity   77.3%; Pred. No. 1.4e-233;
Matches 1474; Conservative 0; Mismatches 415; Indels 18; Gaps 5;

Qy   170 TATCAAACAGCACTGTCACTACTCATGCACAAAACTGCCTCCC---------AAAGACTTTT 223
              |||| ||| ||||||||||||||||||||||||||| |||||||||
Db   640 TATTAAAACTGATAAAAGCAAAAGCCAATGCACAAAAACTACCTCCCTAGAGAAAGGCTAGTC 699

Qy   224 CCCAGGTCCCTCGTATCAAAACATTAAGAGAGTATAAATAGCACAGATCTCTTGTCAGA 283
              |||| |||||||||||||||||||||||||||||||||||| ||||| || ||
Db   700 CCTTTTCTTCCCCATTCATTTCATTTATTGATGACATAGTAGAAAACAGCATATTCTTATCAAA 759

Qy   284 TTGGACAAAACAG---CAACAAACAAAAAATGAAGTATAGAC 340
              ||||||||| ||||||||||||||||||||||
Db   760 TTTGATGAAAAAGCCCCAACACGTTTGAACTGGTGTCATGTGAACTGTACCG 819

Qy   341 AATGTCTACATAATTCAACTTTCCCCGCCGCGGGGTGCCTGTCTCCAGAAAGGAGTCTTGCTCG 400
              |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   820 AAATGTCTACGTAATTCCACTTTTCCTGCTGCGGGGTTCCTGTCYCAGAAGAGAGTCTTTGCTCG 879

Qy   401 TGCTGGTTTTTTATTATACTGGTGTGAATGACAAGGTCTAAAYGCTTCTTCTTGTTGTGGCCTGAT 460
              |||||||| |||||||||||||||||| ||||||||||||||||||||||||||||||||
Db   880 TGCTGGTTTCTATTACACTGGGTGTGAAATGACAAGGTCAAAXGCTTCTGTTGTGGCCCTGAT 939

Qy   461 GCTGGATAAACTGGAGAGACAGAGTCCTATTCAAAAGCAATAAACAGCTATATATCCTAG 520
              |||||||||| |||||| |||| ||||||| ||||||| ||||||||||||
Db   940 GCTGGATAACTGGAAAGGAGACAGCAGTCCTACTGAAAAGXATAAAAAAGTTGTATCCTAG 999

Qy   521 CTGTAGCTTTTATTCAGAAGAATCTGGGTTTTCAGCT----AGTCTGXGGATCXACCTCTAAGAAATAC 577
              |||||| ||| |||||| || |||||||||| ||||||||||| ||||||||||
Db   1000 CTGCAGAATTCGTTCAGAGTTCTAAATTCCGGTTAACGCTACCTCTCAGCCTAC 1059

Qy   578 GT----CTCCAATGAGAAACAGTTTTGCACATTGCACATTCCACCTTGGGAACATAGTAG 634
              || | |||||| ||||||||||||||||||||||||||||
Db   1060 TTTTCCTTCTTCAGTAACAAAATTCCACACTCATTACTTCGGGGTACAGAAAACAGTGG 1119

Qy   635 CTTGTTCAGTGGTTCTTACTCCAGCCTTTCTCCAAACCCTTTAATTCTTAGAGCAGTTGA 694
              ||||||| | ||||||| ||||||||||||||||| ||||||||||||
Db   1120 ATATTTCCGAGGCTCTTATTCCAACACTCTCCAATCCACAAAAACTCGTTTAAACTTCCAGAGCAAAATCA 1179

Qy   695 AGACATCTCTTCAATCGAGGAGCTACACCCTACCAGTTAATGCAATGAGTACTGAAGAAGCCAG 754
              ||| ||| |||| ||||||||||||||| ||| |||||||| ||||||||||
Db   1180 AGATTTTTCTGCCTTGATGAGAAGTTCCTACCACTGTGCAATGAATAACGAAAATGCCAG 1239

Qy   1172 TGCCAAGTGCGGTTTTCCAAGGGTGTGAGTTCTTGATACGAATGAAAAGGCCAAGAGCGTTTGTTGA 1231
              |||| || | |||||||||| ||||| ||||||||| ||||||||||| ||||||||
Db   1660 TGCCAAGTGGGTTTTCCCAAGGTGTGCTGAGTTCATCTTGATAAGGAAATAAAGGACAAGGAGTTCATCCG 1719

Qy   1112 CTTTTGTTGTTGGATGGGTGGCTTGAGGTGTTGGGGAATCTGGGAGATGATCGGAGCAACA 1171
              ||| |||||||||||| |||| |||||||||||||||||| ||||| |||||
Db   1540 TCCTGAGCAGCTTGCAAGTGCGGCGGGGTTTTTATTATGGGGTTAACAGTGATAGTCAAAATG 1599

Qy   1052 GCCTGAGCAGCTTGCCAAGTCGCTGGTTTCTTTTTTATTATGTGGTCGGGTCGGGTCAAATG 1111
              ||||||||| |||||| |||| || |||||||||||||||||||||||| ||
Db   1480 GCAGACACATGCAGCAGCCCGCCTTTAAAACAATTCTTTAAACTGGCCCCTTCTCTAGTTTGA 1539

Qy   992 GCAGACACATGCAGCTCGAAATGAGAACATTTATGTGTACTGGCGGCCATCTAGTTCCAGTTCA 1051
              ||||| |||||| || ||||||||| ||| ||||||||||||||||||||||||||||
Db   1420 CCCATTTATTAGAAAATCAGCTTCAAGACACTTCAAGDATACACAGTTTTCTAATCTGAGCAT 1479

Qy   935 TCCATTTTTGAAAAAT---TCTCTAGAAACTCTCAGAGTTGCATTTCAAAATCTGAGCAT 991
              ||| |||||| |||| ||||| ||||||||||| ||||||||||||||||||
Db   1360 GAGCAATTGGGAAACCGGAAGGATAATGCTATGTCAGAACAACACCTGAGACATTTTCCCAAATG 1419

Qy   815 TGGTTTTTTATTATATAGAGACCTGGAGATAGGGGTAGCCTGCTTTTTGTTGCCAACTG 874
              ||||||||| |||||||||| || |||| | ||||| |||||||||||||| |||
Db   1300 AGGCTTTTTACTTACATAGGACCTGGAACAGAGTGGCTTGGCTTTTCTGTCGGCCAAACAGATCTGGCCAAATG 1359

Qy   755 ATTTTCTTACCTTACCCATATGTGGCCATTAACTTTTTTGTTGTCACCATACAGAATTGGCCAAGAGC 814
              ||||| ||||||||||| ||| |||||||||||||||| |||||| || ||| ||
Db   1240 ATTACTTACTTTTCAGACATGGCCATTGCACTGGGCCAAAGC 1299

Qy   1232 TGAGATTCAAGGTAGATATCCTCATCTTCCTTGACAGCGTGTTGTCAACTTCAGATACCAC 1291
              ||||||||| ||||| ||||||||||||||||||| ||| ||||||||||||
Db   1720 TCAAGTTCAAGGCCAGTTACCCTCATCTACTTGAACAGCTTCCACATCCAATCAGACAGCCC 1779

Qy   1292 TGGAGAAGAAAAATGCTGACCCACCAATTATTCATTTCATTTGATCCTGGAGGAAAGTTCTTCAGA 1351
              ||| |||||||||||||||||||| |||||||||||| ||||||||||||||||| |||
Db   1780 AGGAGAGGAAAAAATGCAGAGTCATCAAATAATTTGAACCTTGGAAGAGACCCATTCAGA 1839

Qy   1352 AGATGCTGTCATGATGAATACACCTGTGGTTAAATCTGCCTTGGAAAGTGGGCTTTAATAG 1411
              ||||||||||||||||| ||||||||| |||||||||| ||||||||| |||||||
Db   1840 AGATGCAATCATGATGATGAATACTCCCTGTGATTAAATGCTGACCGCCTGGAAATGGGCCTTTAGTAG 1899

Qy   1412 AGACCTGGGTGAAACAACCAGAGTTCAAAGTAAAAATCCTGACAACTGGAGGAGAACTATATAAAAC 1471
              |||||||| |||||||| ||| |||||||||| |||||||| |||||||||||||
Db   1900 AAGCCTGGGTAAAAACAGACAGTTCAAAGAAAAATCCTAGCAACTGGCGAGAGAATTATAGACT 1959

Qy   1472 AGTTAATGATATTGTGTCAGCACTTCTAAAATGCTGAAAGATGAAAAGAAAGAGGAGAGGAGA 1531
              ||||| |||| ||||||||| ||||||| ||||| |||||||||| ||||||||||
Db   1960 AGTCAATGATCTTGTTGTTAGCTCAAATGCAGAAGATGAAAATAAGGGAAGAGGAGAG 2019

Qy   1532 GGGAAAACAAGCTGAAGGAAGAAATGGCATCAGATGATGATTTGTCAATTAATTCGGAGAAGAAT 1591
              |||| |||||||||| |||||||||| |||||||||| |||||||||||| ||||
Db   2020 AGAAAGAGCAACTGAGGAAAAAGAAAATCAAAATGATTATTAATTATTCCGGAAGAAATAGAAT 2079

Qy   1592 GGCTCTCTTCAACAATTGACATGTGCTTCCTATCCTGGATAATCTTTTAAAGGGCCAA 1651
              ||| |||||| || |||| |||| || ||| ||||||||||| |||
Db   2080 GGCACTTTTCAAACACTTTGACTTGCTGTAAACTGCGCCGG 2139

Qy   1652 TGTAATTAATTAAAACAGGAACACATGAATATATTTAAACAAAAACACAGATACCTTTACAAGC 1711
              ||||||| |||||| ||||| ||||||||||| || ||||||| ||||||
Db   2140 AATTATTAATGAACAAGAACAATGATGTTAAATCGTATTTATTAAACAGAGAGC 2199

Qy   1712 GAGAGAACTGATTGATACCATTTGGTTAAAGGAAAATGTCGGGCCAAACATCTTCAAAAA 1771
              |||||||||||||||| ||||||||||||| |||||||||| ||||||||||
Db   2200 AAGAGAACTGATTGAATACGATTTAGTAAAAGGAAATATATTGCAAGCACAGTTTGAAAG 2259

Qy   1772 CTGTCTCAAAAGGAAATTGACTCTACATTCTGATAAGAACTTATTGTTGGATAGAATAATGAA 1831
              ||||| ||||||||||||||||| || || ||||||||||| ||||||||||
Db   2260 CTCTCTGCAAGAAGCTGAAGCTGTGTTATATGAGACCATTATTTTGTGCAACAGGACAATAAA 2319

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RESULT 15
US-10-197-290-1

; Sequence 1, Application US/10197290
; Publication No. US20030083300A1
; GENERAL INFORMATION:
; APPLICANT:  C. Frank Bennett
; APPLICANT:  Elizabeth J. Ackermann
; APPLICANT:  Lex M. Cowsert
; TITLE OF INVENTION: ANTISENSE MODULATION OF CELLULAR INHIBITOR OF APOPTOSIS-2
; TITLE OF INVENTION: EXPRESSION
; FILE REFERENCE: RTSP-0421
; CURRENT APPLICATION NUMBER: US/10/197,290
; CURRENT FILING DATE: 2002-07-16
; PRIOR APPLICATION NUMBER: 09/857,299
; PRIOR FILING DATE: 2001-20-04
; PRIOR APPLICATION NUMBER: PCT/US99/22083
; PRIOR FILING DATE: 1999-09-23
; NUMBER OF SEQ ID NOS: 47
; SEQ ID NO 1
; LENGTH: 3076
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (725)..(2539)
US-10-197-290-1

Query Match          44.7%;  Score 1157;  DB 14;  Length 3076;
Best Local Similarity  77.3%;  Pred. No. 1.4e-233;
Matches 1474;  Conservative 0;  Mismatches 415;  Indels 18;  Gaps 5;
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Qy  1832  GTATATTCCAACGAAGAAGTGTTTCAGGTCGTGTCACTGGAGAGAACAATTGAGGAGGGTTGCA  1891
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Db  2320  ATATATTCCCACAGAAGATGTTTCAGAGTCACCAGTGGAAGAACAATTGCGGAGGACTACA    2379

Qy  1892  AGAAGAACGAACTTGTAAAGTGTGTGATGGACAAAGAAGTTTCTGTTGTTGTATTTAATTCCTTG  1951
          |||||||| |||| | |||| || || ||||||||||||| | |||| || |||| ||||||||
Db  2380  AGAAGAAAGAAACATGTGTATGGGACAAAGAAGTGTCCATAGTGTTGTTTAATTCCTTG       2439

Qy  1952  TGGTCATCTGGTAGTAGTGCCAGGAATGTGCCCCTTCTCTAAGAAAAATGCCCTATTTGCAG    2011
          ||||||||| ||||||| || |||| ||||| | |||||||||||||| |||| ||||| |||
Db  2440  TGGTCATCTAGTAGTATGCGAAGATATTGTGCTCCTTCTTTAAGAAAGTGTCCTATTTGTAG    2499

Qy  2012  GGGTATAATCAAGGGTACTGTTCGTTCGTACATTTTCTCTCTTAAAGAAAAA              2058
          | |||| |||||||||| |||||||| ||||||||| | | |||||||| |
Db  2500  GAGTACAATCAAGGGTACAGTTCGTACATTTCTTTCATGAAGAAGAA                  2546

Qy   170  TATCAAACAGCACTGTCACCTACTCATGCACAAAACTGCCTCCC--------AAAGACTTTT     223
          ||| |||| |||| |||| |||  | ||||||||||||||||        |||||||||
Db   640  TATTAAACTGATAAAAAGACAAAGCCATGCACAAAACTACCTCCCTAGAGAAAGGCTAGTC     699

Qy   224  CCCAGGTCCCTCGTTATCAAAACATTAAGAGAGTATAATGGAAGATAGCACGGATCCTTGTCAGA  283
          ||||||| | ||| | |  ||||| ||  |||||||| |||| | | || ||| || |||||
Db   700  CCTTTTCTCCCCATTCATTCATATGACACATAGTAGAAACAGACATATTCTTATCAAA        759

Qy   284  TTGGACAAAACAG---CAACAAACAAAAAATGAAGTATGACTTTTCCTGTGAACTCTACAG     340
          |||||||| |||   ||| ||| || |||||| ||  ||||||||| |||| |||| ||||
Db   760  TTTGATGAAAAAGCCCAACAACCGTTTGAACTGAAAATACGACTTGTCATGTGAACTGTACCG   819

Qy   341  AATGTCTACATATTCAACTTCCCCGCCGGGGCCTGTCTCTCAGAAGAAGGAGTCTTGCTCG     400
          |||||| ||| | | |||||| || | ||| || || | ||| |||| |||||| |||| |
Db   820  AATGTCTACAACTTTCACCTTTTCCTGCTCTGGGGTTCCTGTGTTCAGGAGTCTTCGCTG      879

Qy   401  TGCTCGGTTTTTATTTATACTCGGTGTGGTGAATGACAAAGGTCAAAATGCT             460
          ||||| |||||| |||||| | ||| | || || || ||||||| |||| |||
Db   880  TGCTGGTTTCTATTTCACTCTTTTCTCGTTGTGTTGGTGCCCTGAT                  939

Qy   461  GCTTGGGAATAACTGGAAACTAGGAGGACAGTCCTATTCAAAAAGCATAAAACAGCTAATATCTCAG  520
          | |||||| | ||| ||  | ||| |||| | ||| | ||||| | | ||||||||| | |||||
Db   940  GCTGGGATAACTGAAAAGAGGAGGACAGTCCTACTGAAAAGCATAAAATGTTGTATCCTTAG    999

Qy   521  CTGTAGCTTTATTCAGAATCTGGGTTTCAGCT---AGTCTGGGATCCACACCTCTCTAGAATAC   577
          ||||| || ||||||| | || || ||||    ||||| ||||| | || || | ||||||
Db  1000  CTGCAGGATTCGTTCAGAGTCTAGAATTCGTTTAGCGATTGCACATTCATTATCTCAGCCTAC  1059
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Qy  1000  CTGGCAGATTCTGTTCAGAGTCTGGAGTTCTAAATTCGTTAACGAACTTGGAAGCTACCTTCTCAGCCTAC  1059
          |||| ||| ||||| ||||| ||| ||| || | |||||  |||  |||| |||| ||  |||||| |||
Db   578  GT---CTCCAATGAGAAACACAGTTTTGCACATTCATTATCTCCGGGTACAACCTTGGAAACAGTGG       634

Qy  1060  TTTTCCTTCTTCAGTAACGAAATTCCACACACTCATTACTTCCGGGTACAGAAAACAGTGG  1119
          |||| |||| |||||||| ||| ||||| |||||||| |||||  |||| ||  |||
Db  1060  TTTTCCTTCTTCAGTAACGAAATTCCACACACTCATTACTTCCGGGTACAGAAAACAGTGG  1119

Qy   635  CTTGTTCAGTGGGTTCTTACTCCAGCCCTTTCTCCAAACCCTTCTTAATTCTAGAGCAGTTGA     694
          |||| ||| ||||| ||||| || ||  | || |||| || ||||||||||||| |||| |
Db  1120  ATATTTCCGTGGCTCTTATTCAAAACTCTCCATCAAAATCCTGTAAAATCCTAGAGCAAATCA  1179

Qy   695  AGACAATCTCTTCATCGAGGACTAACCCCTACAGTTTATGCAATGAGTACTGAAGAGCCAG     754
          |||| | | | |||| ||||| |||||| ||||||| | |||||| ||||||||  |||||
Db  1180  AGATTTTTCTGCCCTTGATGAGAAGTTCCTACCACTGTGCAATGAATAACGAAAAATGCCAG    1239

Qy   755  ATTTCTTACCTACCATATGTGGCCATTAAACTTTTTTGTGCACCATCAGAATTGGCAAGAGC     814
          |||||||| || || ||||||||||| |||||| | || ||| |||||||| || |||| |
Db  1240  ATTACTTACTTTTCAGACATGGCCATTGGACTTTTCTGTCGCAACAGATCTGGACAATGGC    1299

Qy   875  CAGTAACTGGGAACCAAAAGGATGAGTGCTATGTCAGAAACACCGGAGGCCATTTCCCAACTG   934
          ||||  |||||||| | |||| |||| ||||| ||||| ||||  ||  ||| || |  ||
Db  1300  AGGCGTTTTACTACATTAGGACACCGTGGGGTAGCCTTGTCTGCTTTGCCTGTGGTGGGAAAAT  1359

Qy   935  TCCATTTTTGGAAAAT---TCTCTAGAAAACTCTGAGGTTTGCATTCAAAATCTGAGCAT    991
          |||| | |||||||    | | ||| |||| || || ||| |||| || | || |||
Db  1360  GAGCAAATTGGGAACCGAACGGAATAAATGCTAATGTCAGAACACCTGAGAACATTTCCCAATG  1419

Qy   992  GCAAGACACATGCGAGCTCGAATGAGAACATTTATGTGACTGGAGCCATCTAGTGTTCCAGTCA  1051
          |||  | | ||| ||| | ||| || ||| | |||  | || || || | |  |||| |||
Db  1420  CCCATTTATAGAAAATCAGCTTCAAGACACTTCAAGATACAGTTTCTAAATCTGAGCAT      1479

Qy  1052  GCCTGAGCAGCTTGCAAGGTGCTGGTTTTTTATTATTGGTTGCGCAATGATGATGTCAAATG  1111
          |||  |||||||| ||| |||  | ||| |||| |  |||| |  |||||||| || ||||
Db  1480  GCAGACACATGCAGCCCGGCTTTAAAACATTCTTTAAACTGGCCCCTCTAGTTGTCTAGTTAA   1539

Qy  1112  CTTTTGTTGTGATGGTGGTGGCTTGAGGTGTTGGGAAATCTGGAGATGATCCATGGGTAGAACA  1171
          ||||||| || |||| | | || ||| |||| |||||||||||||||| |  |||
Db  1540  TCCTGAGCAGCTTGCAAGTGGCGGGTTTTTTATTATGGTGGGAACAGTGAATGTCAAATG     1599

Qy  1172  TGCCAAGTGGTTTCCAAGGTGTGAGTTCTTGATACGAATGAAAAGGCCAAGAGTTTGTTGA  1231
          ||||||||| ||| ||||| ||||| | || ||| || | |||| |||||||| |||||
Db  1600  CTTTTGCTTGTGATGGTGGACTCAGGTGTTGGGAATCTGGAGATGATCCATGGGTTCAACA  1659

Qy  1232  TGAGATTCAAGGTAGATATCCTCATCTTCTTGAACAGCGTTGTCAACTTCAGATACCAC    1291
          ||||| ||||| || ||| | | | || ||| ||||  ||||| ||||| |||| ||||
Db  1720  TCAAGTTCAAGGCCAGTTACCCTCATCACTCTACTTGAACAGCTCCACCATCAGACAGCCC    1779

Qy  1292  TGGGAGAGAAAATGCTGACCCACCAATATTCATCTTTGACCTGAGAGAAAGTTCTTCAGA  1351
          ||||||||||||| || ||   || |||| ||| ||||| ||||||| || ||||| ||
Db  1780  AGGGAGATGAAAATGCAGAGTCATCAATTATCCATTTGAACCTGGAGAAGACCATTCAGA    1839

Qy  1352  AGATGCTGTCATGATGAATAACACCTGTGGTTAAATCTGCCTTGGAAAATGGGGCTTTAATAG  1411
          |||| | |  || ||||||| ||| || |||  |||  | ||| |||||||| ||| ||||
Db  1900  AAGCCTGGGTAAAACAGTTCAAAAGTAAAAATCCTGACAAACTGGAGACTATAGACT        1959

Qy  1412  AGACCTGGTGAAACAAAACAGTTCAAAAGTAAAAATCCTGACAACTGGAGAACTATAAAAAC  1471
          || || ||| |||| |||||||||||| |||||| ||||||| |  ||| |  ||||
Db  1840  AGATGCAAATCAGTGAATAAGTCCCTGTGATTAATGAAAATGGGGCTTTAGTAG           1899

Qy  1472  AGTTAATGATATGTCTGCATCAGATGAGTCCAACTTCTAAAATGTGAAGATGAAGGAGGAGA  1531
          || |||| | || |||||||||||| || |||| ||||| ||||| |||||||| ||| |
Db  1960  AGTCAATGATCTTGTGTTTAGACTTACTCGAAGAGATGAAATAAGGAAGGAGAGGAG        2019

Qy  1532  GGAAAAACAAGCTGAAGAAATTGCATCAGATGATTGTTGTCATTATTGTCTTAAGATAGAAT  1591
          |||| || ||||||| ||| || |  |||| ||||| ||| ||| | ||||| | |
Db  2020  AGAAAGAGCAACTGAGGAAAAAGAATCAAATGATTTATTATTTAATCCGGAAGAATATGAAT    2079

Qy  1592  GGCTCTCTTTCAACAATTGACATGGTGTGCTTCCTATCCTGGATAATCTTTTAAAGGCCAA  1651
          |||| ||||||||| | || | || || || ||| |||  || ||  |  || ||| |||
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Db  2080 GGCACTTTTTCAACATTTGACTTGTGTGTAATTCCAATCCTGGATAGTCTACTAACTGCCGG 2139
              ||||||   |||||| |||||||||||||||||||
Qy  1652 TGTAATTAATAAACAGGAACATGATATTATTATTAAACAAAAAACAGATACCTTTACAAGC 1711
              ||||||   |||||| |||||||||||||||||
Db  2140 AATTATTAATGAACAAGAACATGATGTTATTATTAAACAGAAGAGCACAGAGACGTCTTTACAAGC 2199

Qy  1712 GAGAGAACTGATTGATTACCATTTTTGGTTAAAGGAAATGCTTGCGGGCCAACATCTTCAAAAA 1771
              ||||||   |||||| ||||||   |||||
Db  2200 AAGAGAACTGATTGATTACGATTTTTAGTAAAAGGGAAATATTTGCAGCCACTGTTATTCAGAAA 2259

Qy  1772 CTGTCTAAAAGAAATTGACTCTACATTGTATAAGAACTTATTTGTGGATAAGAAATATGAA 1831
              ||||||   |||||| |||||||||||||||
Db  2260 CTCTCTTGCAAGAAGCTGAAGCTGTGGTTATATGAGCATTTATTTGTGCAACAGCAGGACATAAA 2319

Qy  1832 GTATATTCCAACAGAAGATGTTTCAGGTCTCGTCACTGGAAGAACAATTGAGGAGGTTGCA 1891
              ||||||   |||||| |||||||||||||||
Db  2320 ATATATTCCCACAGAAGATGTTTCAGATCTACCAGTGGAAGAACAATTGCGGAGACTACA 2379

Qy  1892 AGAAGAACGAACTTGTAAAGTGTGTATGGACAAAGAAGTTTCTGTTGTTATTTATTCCTTG 1951
              ||||||   |||||| |||||||||||||||
Db  2380 AGAAGAAAGAACATGTAAAAGTGTGTATGGACAAAGAAGTGTCCATAGTGTTTTATTCCTTG 2439

Qy  1952 TGGTCATCTGGTAGTATGCCAGGAAATGTGCCCCTTCTCTAAGAAAAATGCCCTATTTGCAG 2011
              ||||||   |||||| |||||||||||||||
Db  2440 TGGTCATCTAGTAGTATGCCAAAGATTGTGCTCCCTTCTTCTTTTAAGAAAAGTTGTCCTATTTGTAG 2499

Qy  2012 GGGTATAATCAAGGGTACTGTTTCGTACATTTTCTCTCTTAAAGAAAAA 2058
              ||||||   |||||| |||||||||||||||
Db  2500 GAGTACAAATCAAGGGTACAGTTCGTTACACATTTCTTTCATTGAAGAAGAA 2546
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